



SEQUENCE LISTING

<110> LEY, Arthur C.
GUTERMAN, Sonia K.
MARKLAND, William
KENT, Rachel B.
ROBERTS, Bruce L.
LADNER, Robert C.

<120> ITI-D1 KUNITZ DOMAIN MUTANTS AS HNE INHIBITORS

<130> D0617.7005US01

<140> 10/038,722

<141> 2002-01-08

<150> US 08/849,406

<151> 1999-07-21

<150> PCT/US95/16349

<151> 1995-12-15

<150> US 08/358,160

<151> 1994-12-16

<150> US 08/133,031

<151> 1992-02-28

<160> 140

<170> PatentIn version 3.1

<210> 1

<211> 276

<212> DNA

<213> Artificial Sequence

<220>

<223> IIIsp::bpti::matureIII (initial fragment)

<400> 1

gtgaaaaaat tattattcgc aattccttta gttgttcctt tctattctgg cgcccgcccg 60

gatttctgtc tcgagccacc atacactggg ccctgcaaag cgcgcatcat ccgctatttc 120

tacaatgcta aagcaggcct gtgccagacc tttgtatacg gtgggtgccg tgctaagcgt 180

aacaacttta aatcggccga agattgcatt cgctacctgc gtggcgccgc tgaaactgtt 240

gaaagttgtt tagcaaaacc ccatacagaa aattca 276

<210> 2

<211> 92

<212> PRT

<213> Artificial Sequence

<220>

<223> IIIsp::bpti::matureIII (initial fragment)

<400> 2

Met Lys Lys Leu Leu Phe Ala Ile Pro Leu Val Val Pro Phe Tyr Ser
1 5 10 15
Gly Ala Arg Pro Asp Phe Cys Leu Glu Pro Pro Tyr Thr Gly Pro Cys
20 25 30
Lys Ala Arg Ile Ile Arg Tyr Phe Tyr Asn Ala Lys Ala Gly Leu Cys
35 40 45
Gln Thr Phe Val Tyr Gly Gly Cys Arg Ala Lys Arg Asn Asn Phe Lys
50 55 60
Ser Ala Glu Asp Cys Met Arg Thr Cys Gly Gly Ala Ala Glu Thr Val
65 70 75 80
Glu Ser Cys Leu Ala Lys Pro His Thr Glu Asn Ser
85 90

<210> 3

<211> 285

<212> DNA

<213> Artificial Sequence

<220>

<223> IIIsp::itiD1::mature III fusion gene

<400> 3

gtgaaaaaat tattattcgc aattccttta gttgttcctt tctattctgg cgccaaagaa 60
gactcttgcc agctgggcta ctcggccggt ccctgcatgg gaatgaccag caggtatttc 120
tataatggta catccatggc ctgtgagact ttccagtaag gcggtgcat gggcaacggt 180
aacaacttcg tcacagaaaa ggagtgtctg cagacctgcc gaactgtggg cgccgctgaa 240
actgttgaaa gttgtttagc aaaaccccat acagaaaatt cattt 285

<210> 4

<211> 95

<212> PRT

<213> Artificial Sequence

<220>

<223> IIIsp::itiD1::mature III fusion gene

<400> 4

Met Lys Lys Leu Leu Phe Ala Ile Pro Leu Val Val Pro Phe Tyr Ser
1 5 10 15

Gly Ala Lys Glu Asp Ser Cys Gln Leu Gly Tyr Ser Ala Gly Pro Cys
20 25 30

Met Gly Met Thr Ser Arg Tyr Phe Tyr Asn Gly Thr Ser Met Ala Cys
35 40 45

Glu Thr Phe Gln Tyr Gly Gly Cys Met Gly Asn Gly Asn Asn Phe Val
50 55 60

Thr Glu Lys Glu Cys Leu Gln Thr Cys Arg Thr Val Gly Ala Ala Glu
65 70 75 80

Thr Val Glu Ser Cys Leu Ala Lys Pro His Thr Glu Asn Ser Phe
85 90 95

<210> 5
<211> 58
<212> PRT
<213> Artificial Sequence

<220>

<223> Consensus Kunitz domain

<400> 5

Arg Pro Asp Phe Cys Leu Leu Pro Ala Glu Thr Gly Pro Cys Arg Ala
1 5 10 15

Met Ile Pro Arg Phe Tyr Tyr Asn Ala Lys Ser Gly Lys Cys Glu Pro
20 25 30

Phe Ile Tyr Gly Gly Cys Gly Gly Asn Ala Asn Asn Phe Lys Thr Glu
35 40 45

Glu Glu Cys Arg Arg Thr Cys Gly Gly Ala
50 55

<210> 6
<211> 58
<212> PRT
<213> Bos Taurus

<400> 6

Arg Pro Asp Phe Cys Leu Glu Pro Pro Tyr Thr Gly Pro Cys Lys Ala
1 5 10 15

Arg Ile Ile Arg Tyr Phe Tyr Asn Ala Lys Ala Gly Leu Cys Gln Thr
20 25 30

Phe Val Tyr Gly Gly Cys Arg Ala Lys Arg Asn Asn Phe Lys Ser Ala
35 40 45

Glu Asp Cys Met Arg Thr Cys Gly Gly Ala
50 55

<210> 7
<211> 58
<212> PRT
<213> Artificial Sequence

<220>

<223> Epi-HNE-1

<400> 7

Arg Pro Asp Phe Cys Leu Glu Pro Pro Tyr Thr Gly Pro Cys Ile Ala
1 5 10 15
Phe Phe Pro Arg Tyr Phe Tyr Asn Ala Lys Ala Gly Leu Cys Gln Thr
20 25 30
Phe Val Tyr Gly Gly Cys Met Gly Asn Gly Asn Asn Phe Lys Ser Ala
35 40 45
Glu Asp Cys Met Arg Thr Cys Gly Gly Ala
50 55

<210> 8
<211> 62
<212> PRT
<213> Artificial Sequence

<220>

<223> Epi-HNE-2

<400> 8

Glu Ala Glu Ala Arg Pro Asp Phe Cys Leu Glu Pro Pro Tyr Thr Gly
1 5 10 15
Pro Cys Ile Ala Phe Phe Pro Arg Tyr Phe Tyr Asn Ala Lys Ala Gly
20 25 30
Leu Cys Gln Thr Phe Val Tyr Gly Gly Cys Met Gly Asn Gly Asn Asn
35 40 45
Phe Lys Ser Ala Glu Asp Cys Met Arg Thr Cys Gly Gly Ala
50 55 60

<210> 9
<211> 58
<212> PRT
<213> Artificial Sequence

<220>

<223> EpiNE7

<400> 9

Arg Pro Asp Phe Cys Leu Glu Pro Pro Tyr Thr Gly Pro Cys Val Ala
1 5 10 15

Met Phe Pro Arg Tyr Phe Tyr Asn Ala Lys Ala Gly Leu Cys Gln Thr
20 25 30

Phe Val Tyr Gly Gly Cys Met Gly Asn Gly Asn Asn Phe Lys Ser Ala
35 40 45

Glu Asp Cys Met Arg Thr Cys Gly Gly Ala
50 55

<210> 10
<211> 58
<212> PRT
<213> Artificial Sequence

<220>

<223> EpiNE3

<400> 10

Arg Pro Asp Phe Cys Leu Glu Pro Pro Tyr Thr Gly Pro Cys Val Gly
1 5 10 15

Phe Phe Ser Arg Tyr Phe Tyr Asn Ala Lys Ala Gly Leu Cys Gln Thr
20 25 30

Phe Val Tyr Gly Gly Cys Met Gly Asn Gly Asn Asn Phe Lys Ser Ala
35 40 45

Glu Asp Cys Met Arg Thr Cys Gly Gly Ala
50 55

<210> 11
<211> 58
<212> PRT
<213> Artificial Sequence

<220>

<223> EpiNE6

<400> 11

Arg Pro Asp Phe Cys Leu Glu Pro Pro Tyr Thr Gly Pro Cys Val Gly
1 5 10 15

Phe Phe Gln Arg Tyr Phe Tyr Asn Ala Lys Ala Gly Leu Cys Gln Thr
20 25 30

Phe Val Tyr Gly Gly Cys Met Gly Asn Gly Asn Asn Phe Lys Ser Ala
35 40 45

Glu Asp Cys Met Arg Thr Cys Gly Gly Ala
50 55

<210> 12
<211> 58
<212> PRT

<213> Artificial Sequence

<220>

<223> EpiNE4

<400> 12

Arg Pro Asp Phe Cys Leu Glu Pro Pro Tyr Thr Gly Pro Cys Val Ala
1 5 10 15

Ile Phe Pro Arg Tyr Phe Tyr Asn Ala Lys Ala Gly Leu Cys Gln Thr
20 25 30

Phe Val Tyr Gly Gly Cys Met Gly Asn Gly Asn Asn Phe Lys Ser Ala
35 40 45

Glu Asp Cys Met Arg Thr Cys Gly Gly Ala
50 55

<210> 13

<211> 58

<212> PRT

<213> Artificial Sequence

<220>

<223> EpiNE8

<400> 13

Arg Pro Asp Phe Cys Leu Glu Pro Pro Tyr Thr Gly Pro Cys Val Ala
1 5 10 15

Phe Phe Lys Arg Ser Phe Tyr Asn Ala Lys Ala Gly Leu Cys Gln Thr
20 25 30

Phe Val Tyr Gly Gly Cys Met Gly Asn Gly Asn Asn Phe Lys Ser Ala
35 40 45

Glu Asp Cys Met Arg Thr Cys Gly Gly Ala
50 55

<210> 14

<211> 58

<212> PRT

<213> Artificial Sequence

<220>

<223> EpiNE5

<400> 14

Arg Pro Asp Phe Cys Leu Glu Pro Pro Tyr Thr Gly Pro Cys Ile Ala
1 5 10 15

Phe Phe Gln Arg Tyr Phe Tyr Asn Ala Lys Ala Gly Leu Cys Gln Thr
20 25 30

Phe Val Tyr Gly Gly Cys Met Gly Asn Gly Asn Asn Phe Lys Ser Ala
35 40 45

Glu Asp Cys Met Arg Thr Cys Gly Gly Ala
50 55

<210> 15
<211> 58
<212> PRT
<213> Artificial Sequence

<220>

<223> EpiNE2

<400> 15

Arg Pro Asp Phe Cys Leu Glu Pro Pro Tyr Thr Gly Pro Cys Ile Ala
1 5 10 15

Leu Phe Lys Arg Tyr Phe Tyr Asn Ala Lys Ala Gly Leu Cys Gln Thr
20 25 30

Phe Val Tyr Gly Gly Cys Met Gly Asn Gly Asn Asn Phe Lys Ser Ala
35 40 45

Glu Asp Cys Met Arg Thr Cys Gly Gly Ala
50 55

<210> 16
<211> 58
<212> PRT
<213> Homo sapiens

<400> 16

Lys Glu Asp Ser Cys Gln Leu Gly Tyr Ser Ala Gly Pro Cys Met Gly
1 5 10 15

Met Thr Ser Arg Tyr Phe Tyr Asn Gly Thr Ser Met Ala Cys Glu Thr
20 25 30

Phe Gln Tyr Gly Gly Cys Met Gly Asn Gly Asn Asn Phe Val Thr Glu
35 40 45

Lys Asp Cys Leu Gln Thr Cys Arg Thr Val
50 55

<210> 17
<211> 58
<212> PRT
<213> Artificial Sequence

<220>

<223> BITI-E7-141

<400> 17

Arg Pro Asp Phe Cys Gln Leu Gly Tyr Ser Ala Gly Pro Cys Val Ala
1 5 10 15
Met Phe Pro Arg Tyr Phe Tyr Asn Gly Thr Ser Met Ala Cys Gln Thr
20 25 30
Phe Val Tyr Gly Gly Cys Met Gly Asn Gly Asn Asn Phe Val Thr Glu
35 40 45
Lys Asp Cys Leu Gln Thr Cys Arg Gly Ala
50 55

<210> 18

<211> 58

<212> PRT

<213> Artificial Sequence

<220>

<223> MUTT26A

<400> 18

Arg Pro Asp Phe Cys Gln Leu Gly Tyr Ser Ala Gly Pro Cys Val Ala
1 5 10 15
Met Phe Pro Arg Tyr Phe Tyr Asn Gly Ala Ser Met Ala Cys Gln Thr
20 25 30
Phe Val Tyr Gly Gly Cys Met Gly Asn Gly Asn Asn Phe Val Thr Glu
35 40 45
Lys Asp Cys Leu Gln Thr Cys Arg Gly Ala
50 55

<210> 19

<211> 58

<212> PRT

<213> Artificial Sequence

<220>

<223> MUTQE

<400> 19

Arg Pro Asp Phe Cys Gln Leu Gly Tyr Ser Ala Gly Pro Cys Val Ala
1 5 10 15
Met Phe Pro Arg Tyr Phe Tyr Asn Gly Thr Ser Met Ala Cys Glu Thr
20 25 30
Phe Val Tyr Gly Gly Cys Met Gly Asn Gly Asn Asn Phe Val Thr Glu
35 40 45
Lys Asp Cys Leu Gln Thr Cys Arg Gly Ala
50 55

<210> 20
<211> 58
<212> PRT
<213> Artificial Sequence

<220>

<223> MUT1619

<400> 20

Arg Pro Asp Phe Cys Gln Leu Gly Tyr Ser Ala Gly Pro Cys Val Gly
1 5 10 15
Met Phe Ser Arg Tyr Phe Tyr Asn Gly Thr Ser Met Ala Cys Gln Thr
20 25 30
Phe Val Tyr Gly Gly Cys Met Gly Asn Gly Asn Asn Phe Val Thr Glu
35 40 45
Lys Asp Cys Leu Gln Thr Cys Arg Gly Ala
50 55

<210> 21
<211> 58
<212> PRT
<213> Artificial Sequence

<220>

<223> ITI-D1E7

<400> 21

Lys Glu Asp Ser Cys Gln Leu Gly Tyr Ser Ala Gly Pro Cys Val Ala
1 5 10 15
Met Phe Pro Arg Tyr Phe Tyr Asn Gly Thr Ser Met Ala Cys Glu Thr
20 25 30
Phe Gln Tyr Gly Gly Cys Met Gly Asn Gly Asn Asn Phe Val Thr Glu
35 40 45
Lys Asp Cys Leu Gln Thr Cys Arg Gly Ala
50 55

<210> 22
<211> 58
<212> PRT
<213> Artificial Sequence

<220>

<223> AMINO1

<400> 22

Lys Glu Asp Phe Cys Gln Leu Gly Tyr Ser Ala Gly Pro Cys Val Ala
1 5 10 15

Met Phe Pro Arg Tyr Phe Tyr Asn Gly Thr Ser Met Ala Cys Glu Thr
 20 25 30
 Phe Gln Tyr Gly Gly Cys Met Gly Asn Gly Asn Asn Phe Val Thr Glu
 35 40 45
 Lys Asp Cys Leu Gln Thr Cys Arg Gly Ala
 50 55

<210> 23
 <211> 58
 <212> PRT
 <213> Artificial Sequence

<220>

<223> AMINO2

<400> 23

Lys Pro Asp Ser Cys Gln Leu Gly Tyr Ser Ala Gly Pro Cys Val Ala
 1 5 10 15
 Met Phe Pro Arg Tyr Phe Tyr Asn Gly Thr Ser Met Ala Cys Glu Thr
 20 25 30
 Phe Gln Tyr Gly Gly Cys Met Gly Asn Gly Asn Asn Phe Val Thr Glu
 35 40 45
 Lys Asp Cys Leu Gln Thr Cys Arg Gly Ala
 50 55

<210> 24
 <211> 58
 <212> PRT
 <213> Artificial Sequence

<220>

<223> MUTP1

<400> 24

Arg Pro Asp Phe Cys Gln Leu Gly Tyr Ser Ala Gly Pro Cys Ile Gly
 1 5 10 15
 Met Phe Ser Arg Tyr Phe Tyr Asn Gly Thr Ser Met Ala Cys Glu Thr
 20 25 30
 Phe Gln Tyr Gly Gly Cys Met Gly Asn Gly Asn Asn Phe Val Thr Glu
 35 40 45
 Lys Asp Cys Leu Gln Thr Cys Arg Gly Ala
 50 55

<210> 25
 <211> 58
 <212> PRT

<213> Homo sapiens

<400> 25

Thr Val Ala Ala Cys Asn Leu Pro Ile Val Arg Gly Pro Cys Arg Ala
1 5 10 15
Phe Ile Gln Leu Trp Ala Phe Asp Ala Val Lys Gly Lys Cys Val Leu
20 25 30
Phe Pro Tyr Gly Gly Cys Gln Gly Asn Gly Asn Lys Phe Tyr Ser Glu
35 40 45
Lys Glu Cys Arg Glu Tyr Cys Gly Val Pro
50 55

<210> 26

<211> 56

<212> PRT

<213> Artificial Sequence

<220>

<223> Epi-HNE-3

<400> 26

Ala Ala Cys Asn Leu Pro Ile Val Arg Gly Pro Cys Ile Ala Phe Phe
1 5 10 15
Pro Arg Trp Ala Phe Asp Ala Val Lys Gly Lys Cys Val Leu Phe Pro
20 25 30
Tyr Gly Gly Cys Gln Gly Asn Gly Asn Lys Phe Tyr Ser Glu Lys Glu
35 40 45
Cys Arg Glu Tyr Cys Gly Val Pro
50 55

<210> 27

<211> 56

<212> PRT

<213> Artificial Sequence

<220>

<223> Epi-HNE-4

<400> 27

Glu Ala Cys Asn Leu Pro Ile Val Arg Gly Pro Cys Ile Ala Phe Phe
1 5 10 15
Pro Arg Trp Ala Phe Asp Ala Val Lys Gly Lys Cys Val Leu Phe Pro
20 25 30
Tyr Gly Gly Cys Gln Gly Asn Gly Asn Lys Phe Tyr Ser Glu Lys Glu
35 40 45

Cys Arg Glu Tyr Cys Gly Val Pro
50 55

<210> 28
<211> 58
<212> PRT
<213> Homo sapiens

<400> 28

Val Arg Glu Val Cys Ser Glu Gln Ala Glu Thr Gly Pro Cys Arg Ala
1 5 10 15
Met Ile Ser Arg Trp Tyr Phe Asp Val Thr Glu Gly Lys Cys Ala Pro
20 25 30
Phe Phe Tyr Gly Gly Cys Gly Gly Asn Arg Asn Asn Phe Asp Thr Glu
35 40 45
Glu Tyr Cys Met Ala Val Cys Gly Ser Ala
50 55

<210> 29
<211> 58
<212> PRT
<213> Artificial Sequence

<220>

<223> DPI.1.1

<400> 29

Val Arg Glu Val Cys Ser Glu Gln Ala Tyr Thr Gly Pro Cys Ile Ala
1 5 10 15
Phe Phe Pro Arg Tyr Tyr Phe Asp Val Thr Glu Gly Lys Cys Gln Thr
20 25 30
Phe Val Tyr Gly Gly Cys Met Gly Asn Gly Asn Asn Phe Asp Thr Glu
35 40 45
Glu Tyr Cys Met Ala Val Cys Gly Ser Ala
50 55

<210> 30
<211> 58
<212> PRT
<213> Artificial Sequence

<220>

<223> DPI.1.2

<400> 30

Val Arg Glu Val Cys Ser Glu Gln Ala Glu Thr Gly Pro Cys Ile Ala
1 5 10 15

Met Phe Ser Arg Trp Tyr Phe Asp Val Thr Glu Gly Lys Cys Ala Pro
20 25 30

Phe Val Tyr Gly Gly Cys Gly Gly Asn Arg Asn Asn Phe Asp Thr Glu
35 40 45

Glu Tyr Cys Met Ala Val Cys Gly Ser Ala
50 55

<210> 31

<211> 58

<212> PRT

<213> Artificial Sequence

<220>

<223> DPI.1.3

<400> 31

Val Arg Glu Val Cys Ser Glu Gln Ala Glu Thr Gly Pro Cys Ile Ala
1 5 10 15

Phe Phe Ser Arg Trp Tyr Phe Asp Val Thr Glu Gly Lys Cys Ala Thr
20 25 30

Phe Val Tyr Gly Gly Cys Met Gly Asn Arg Asn Asn Phe Asp Thr Glu
35 40 45

Glu Tyr Cys Met Ala Val Cys Gly Ser Ala
50 55

<210> 32

<211> 58

<212> PRT

<213> Homo sapiens

<400> 32

Asn Ala Glu Ile Cys Leu Leu Pro Leu Asp Tyr Gly Pro Cys Arg Ala
1 5 10 15

Leu Leu Leu Arg Tyr Tyr Tyr Asp Arg Tyr Thr Gln Ser Cys Arg Gln
20 25 30

Phe Leu Tyr Gly Gly Cys Glu Gly Asn Ala Asn Asn Phe Tyr Thr Trp
35 40 45

Glu Ala Cys Asp Asp Ala Cys Trp Arg Ile
50 55

<210> 33

<211> 58

<212> PRT

<213> Artificial Sequence

<220>

<223> DPI.2.1

<400> 33

Asn Ala Glu Ile Cys Leu Leu Pro Leu Tyr Thr Gly Pro Cys Ile Ala
1 5 10 15
Phe Phe Pro Arg Tyr Tyr Tyr Asp Arg Tyr Thr Gln Ser Cys Gln Thr
20 25 30
Phe Val Tyr Gly Gly Cys Met Gly Asn Ala Asn Asn Phe Tyr Thr Trp
35 40 45
Glu Ala Cys Asp Asp Ala Cys Trp Arg Ile
50 55

<210> 34

<211> 58

<212> PRT

<213> Artificial Sequence

<220>

<223> DPI.2.2

<400> 34

Asn Ala Glu Ile Cys Leu Leu Pro Leu Asp Tyr Gly Pro Cys Ile Ala
1 5 10 15
Leu Phe Leu Arg Tyr Tyr Tyr Asp Arg Tyr Thr Gln Ser Cys Arg Gln
20 25 30
Phe Val Tyr Gly Gly Cys Glu Gly Asn Ala Asn Asn Phe Tyr Thr Trp
35 40 45
Glu Ala Cys Asp Asp Ala Cys Trp Arg Ile
50 55

<210> 35

<211> 58

<212> PRT

<213> Artificial Sequence

<220>

<223> DPI.2.3

<400> 35

Asn Ala Glu Ile Cys Leu Leu Pro Leu Asp Thr Gly Pro Cys Ile Ala
1 5 10 15
Phe Phe Leu Arg Tyr Tyr Tyr Asp Arg Tyr Thr Gln Ser Cys Gln Thr
20 25 30
Phe Val Tyr Gly Gly Cys Met Gly Asn Ala Asn Asn Phe Tyr Thr Trp
35 40 45

Glu Ala Cys Asp Asp Ala Cys Trp Arg Ile
50 55

<210> 36
<211> 61
<212> PRT
<213> Homo sapiens

<400> 36

Val Pro Lys Val Cys Arg Leu Gln Val Ser Val Asp Asp Gln Cys Glu
1 5 10 15
Gly Ser Thr Glu Lys Tyr Phe Phe Asn Leu Ser Ser Met Thr Cys Glu
20 25 30
Lys Phe Phe Ser Gly Gly Cys His Arg Asn Arg Ile Glu Asn Arg Phe
35 40 45
Pro Asp Glu Ala Thr Cys Met Gly Phe Cys Ala Pro Lys
50 55 60

<210> 37
<211> 58
<212> PRT
<213> Artificial Sequence

<220>

<223> DPI.3.1

<400> 37

Val Pro Lys Val Cys Arg Leu Gln Val Val Arg Gly Pro Cys Ile Ala
1 5 10 15
Phe Phe Pro Arg Trp Phe Phe Asn Leu Ser Ser Met Thr Cys Val Leu
20 25 30
Phe Pro Tyr Gly Gly Cys Gln Gly Asn Gly Asn Arg Phe Pro Asp Glu
35 40 45
Ala Thr Cys Met Gly Phe Cys Ala Pro Lys
50 55

<210> 38
<211> 61
<212> PRT
<213> Artificial Sequence

<220>

<223> DPI.3.2

<400> 38

Val Pro Lys Val Cys Arg Leu Gln Val Ser Val Asp Asp Gln Cys Ile
1 5 10 15

Gly Ser Phe Glu Lys Tyr Phe Phe Asn Leu Ala Ser Met Thr Cys Glu
 20 25 30
 Thr Phe Val Ser Gly Gly Cys His Arg Asn Arg Ile Glu Asn Arg Phe
 35 40 45
 Pro Asp Glu Ala Thr Cys Met Gly Phe Cys Ala Pro Lys
 50 55 60

<210> 39
 <211> 58
 <212> PRT
 <213> Artificial Sequence

<220>

<223> DPI.3.3

<400> 39

Val Pro Lys Val Cys Arg Leu Gln Val Val Ala Gly Pro Cys Ile Gly
 1 5 10 15
 Phe Phe Lys Arg Tyr Phe Phe Ala Leu Ser Ser Met Thr Cys Glu Thr
 20 25 30
 Phe Val Ser Gly Gly Cys His Arg Asn Arg Asn Arg Phe Pro Asp Glu
 35 40 45
 Ala Thr Cys Met Gly Phe Cys Ala Pro Lys
 50 55

<210> 40
 <211> 58
 <212> PRT
 <213> Homo sapiens

<400> 40

Ile Pro Ser Phe Cys Tyr Ser Pro Lys Asp Glu Gly Leu Cys Ser Ala
 1 5 10 15
 Asn Val Thr Arg Tyr Tyr Phe Asn Pro Arg Tyr Arg Thr Cys Asp Ala
 20 25 30
 Phe Thr Tyr Thr Gly Cys Gly Gly Asn Asp Asn Asn Phe Val Ser Arg
 35 40 45
 Glu Asp Cys Lys Arg Ala Cys Ala Lys Ala
 50 55

<210> 41
 <211> 58
 <212> PRT
 <213> Artificial Sequence

<220>

<223> DPI.4.1

<400> 41

Ile Pro Ser Phe Cys Tyr Ser Pro Lys Ser Ala Gly Pro Cys Val Ala
1 5 10 15
Met Phe Pro Arg Tyr Tyr Phe Asn Pro Arg Tyr Arg Thr Cys Glu Thr
20 25 30
Phe Val Tyr Gly Gly Cys Met Gly Asn Gly Asn Asn Phe Val Ser Arg
35 40 45
Glu Asp Cys Lys Arg Ala Cys Ala Lys Ala
50 55

<210> 42

<211> 58

<212> PRT

<213> Artificial Sequence

<220>

<223> DPI.4.2

<400> 42

Ile Pro Ser Phe Cys Tyr Ser Pro Lys Asp Glu Gly Leu Cys Ile Ala
1 5 10 15
Phe Phe Thr Arg Tyr Tyr Phe Asn Pro Arg Tyr Arg Thr Cys Asp Ala
20 25 30
Phe Thr Tyr Thr Gly Cys Gly Gly Asn Asp Asn Asn Phe Val Ser Arg
35 40 45
Glu Asp Cys Lys Arg Ala Cys Ala Lys Ala
50 55

<210> 43

<211> 58

<212> PRT

<213> Artificial Sequence

<220>

<223> DPI.4.3

<400> 43

Ile Pro Ser Phe Cys Tyr Ser Pro Lys Asp Thr Gly Pro Cys Ile Ala
1 5 10 15
Phe Phe Thr Arg Tyr Tyr Phe Asn Pro Arg Tyr Arg Thr Cys Asp Thr
20 25 30
Phe Val Tyr Gly Gly Cys Gly Gly Asn Asp Asn Asn Phe Val Ser Arg
35 40 45

Glu Asp Cys Lys Arg Ala Cys Ala Lys Ala
50 55

<210> 44
<211> 58
<212> PRT
<213> Homo sapiens

<400> 44

Met His Ser Phe Cys Ala Phe Lys Ala Asp Asp Gly Pro Cys Lys Ala
1 5 10 15

Ile Met Lys Arg Phe Phe Phe Asn Ile Phe Thr Arg Gln Cys Glu Glu
20 25 30

Phe Ile Tyr Gly Gly Cys Glu Gly Asn Gln Asn Arg Phe Glu Ser Leu
35 40 45

Glu Glu Cys Lys Lys Met Cys Thr Arg Asp
50 55

<210> 45
<211> 58
<212> PRT
<213> Artificial Sequence

<220>

<223> DPI.5.1

<400> 45

Met His Ser Phe Cys Ala Phe Lys Ala Ser Ala Gly Pro Cys Val Ala
1 5 10 15

Met Phe Pro Arg Tyr Phe Phe Asn Ile Phe Thr Arg Gln Cys Glu Thr
20 25 30

Phe Val Tyr Gly Gly Cys Met Gly Asn Gly Asn Arg Phe Glu Ser Leu
35 40 45

Glu Glu Cys Lys Lys Met Cys Thr Arg Asp
50 55

<210> 46
<211> 58
<212> PRT
<213> Artificial Sequence

<220>

<223> DPI.5.2

<400> 46

Met His Ser Phe Cys Ala Phe Lys Ala Asp Asp Gly Pro Cys Ile Ala
1 5 10 15

Ile Phe Lys Arg Phe Phe Phe Asn Ile Phe Thr Arg Gln Cys Glu Glu
20 25 30
Phe Ile Tyr Gly Gly Cys Glu Gly Asn Gln Asn Arg Phe Glu Ser Leu
35 40 45
Glu Glu Cys Lys Lys Met Cys Thr Arg Asp
50 55

<210> 47
<211> 58
<212> PRT
<213> Artificial Sequence

<220>

<223> DPI.5.3

<400> 47

Met His Ser Phe Cys Ala Phe Lys Ala Tyr Thr Gly Pro Cys Ile Ala
1 5 10 15
Phe Phe Lys Arg Phe Phe Phe Asn Ile Phe Thr Arg Gln Cys Glu Thr
20 25 30
Phe Ile Tyr Gly Gly Cys Glu Gly Asn Gln Asn Arg Phe Glu Ser Leu
35 40 45
Glu Glu Cys Lys Lys Met Cys Thr Arg Asp
50 55

<210> 48
<211> 58
<212> PRT
<213> Homo sapiens

<400> 48

Lys Pro Asp Phe Cys Phe Leu Glu Glu Asp Pro Gly Ile Cys Arg Gly
1 5 10 15
Tyr Ile Thr Arg Tyr Phe Tyr Asn Asn Gln Thr Lys Gln Cys Glu Arg
20 25 30
Phe Lys Tyr Gly Gly Cys Leu Gly Asn Met Asn Asn Phe Glu Thr Leu
35 40 45
Glu Glu Cys Lys Asn Ile Cys Glu Asp Gly
50 55

<210> 49
<211> 58
<212> PRT
<213> Artificial Sequence

<220>

<223> DPI.6.1

<400> 49

Lys Pro Asp Phe Cys Phe Leu Glu Glu Ser Ala Gly Pro Cys Val Ala
1 5 10 15

Met Phe Pro Arg Tyr Phe Tyr Asn Asn Gln Thr Lys Gln Cys Glu Thr
20 25 30

Phe Val Tyr Gly Gly Cys Met Gly Asn Gly Asn Asn Phe Glu Thr Leu
35 40 45

Glu Glu Cys Lys Asn Ile Cys Glu Asp Gly
50 55

<210> 50

<211> 58

<212> PRT

<213> Artificial Sequence

<220>

<223> DPI.6.2

<400> 50

Lys Pro Asp Phe Cys Phe Leu Glu Glu Asp Pro Gly Ile Cys Val Gly
1 5 10 15

Tyr Phe Thr Arg Tyr Phe Tyr Asn Asn Gln Thr Lys Gln Cys Glu Arg
20 25 30

Phe Lys Tyr Gly Gly Cys Leu Gly Asn Met Asn Asn Phe Glu Thr Leu
35 40 45

Glu Glu Cys Lys Asn Ile Cys Glu Asp Gly
50 55

<210> 51

<211> 58

<212> PRT

<213> Artificial Sequence

<220>

<223> DPI.6.3

<400> 51

Lys Pro Asp Phe Cys Phe Leu Glu Glu Asp Pro Gly Ile Cys Val Gly
1 5 10 15

Phe Phe Thr Arg Tyr Phe Tyr Asn Asn Gln Thr Lys Gln Cys Glu Arg
20 25 30

Phe Val Tyr Gly Gly Cys Leu Gly Asn Met Asn Asn Phe Glu Thr Leu
35 40 45

Glu Glu Cys Lys Asn Ile Cys Glu Asp Gly
50 55

<210> 52
<211> 58
<212> PRT
<213> Artificial Sequence

<220>

<223> DPI.6.4

<400> 52

Lys Pro Asp Phe Cys Phe Leu Glu Glu Asp Pro Gly Ile Cys Val Gly
1 5 10 15
Phe Phe Thr Arg Tyr Phe Tyr Asn Ala Gln Thr Lys Gln Cys Glu Arg
20 25 30
Phe Val Tyr Gly Gly Cys Leu Gly Asn Met Asn Asn Phe Glu Thr Leu
35 40 45
Glu Glu Cys Lys Asn Ile Cys Glu Asp Gly
50 55

<210> 53
<211> 58
<212> PRT
<213> Artificial Sequence

<220>

<223> DPI.6.5

<400> 53

Lys Pro Asp Phe Cys Phe Leu Glu Glu Asp Pro Gly Pro Cys Val Gly
1 5 10 15
Phe Phe Gln Arg Tyr Phe Tyr Asn Ala Gln Thr Lys Gln Cys Glu Arg
20 25 30
Phe Val Tyr Gly Gly Cys Gln Gly Asn Met Asn Asn Phe Glu Thr Leu
35 40 45
Glu Glu Cys Lys Asn Ile Cys Glu Asp Gly
50 55

<210> 54
<211> 58
<212> PRT
<213> Artificial Sequence

<220>

<223> DPI.6.6

<400> 54

Lys Pro Asp Phe Cys Phe Leu Glu Glu Asp Pro Gly Pro Cys Val Gly
1 5 10 15
Phe Phe Thr Arg Tyr Phe Tyr Asn Asn Gln Thr Lys Gln Cys Glu Arg
20 25 30
Phe Val Tyr Gly Gly Cys Gln Gly Asn Met Asn Asn Phe Glu Thr Leu
35 40 45
Glu Glu Cys Lys Asn Ile Cys Glu Asp Gly
50 55

<210> 55

<211> 58

<212> PRT

<213> Artificial Sequence

<220>

<223> DPI.6.7

<400> 55

Lys Pro Asp Phe Cys Phe Leu Glu Glu Asp Pro Gly Pro Cys Ile Gly
1 5 10 15
Phe Phe Pro Arg Tyr Phe Tyr Asn Asn Gln Thr Lys Gln Cys Glu Arg
20 25 30
Phe Val Tyr Gly Gly Cys Gln Gly Asn Met Asn Asn Phe Glu Thr Leu
35 40 45
Glu Glu Cys Lys Asn Ile Cys Glu Asp Gly
50 55

<210> 56

<211> 58

<212> PRT

<213> Homo sapiens

<400> 56

Gly Pro Ser Trp Cys Leu Thr Pro Ala Asp Arg Gly Leu Cys Arg Ala
1 5 10 15
Asn Glu Asn Arg Phe Tyr Tyr Asn Ser Val Ile Gly Lys Cys Arg Pro
20 25 30
Phe Lys Tyr Ser Gly Cys Gly Gly Asn Glu Asn Asn Phe Thr Ser Lys
35 40 45
Gln Glu Cys Leu Arg Ala Cys Lys Lys Gly
50 55

<210> 57

<211> 58

<212> PRT
<213> Artificial Sequence

<220>

<223> DPI.7.1

<400> 57

Gly Pro Ser Trp Cys Leu Thr Pro Ala Val Arg Gly Pro Cys Ile Ala
1 5 10 15
Phe Phe Pro Arg Trp Tyr Tyr Asn Ser Val Ile Gly Lys Cys Val Leu
20 25 30
Phe Pro Tyr Gly Gly Cys Gln Gly Asn Gly Asn Asn Phe Thr Ser Lys
35 40 45
Gln Glu Cys Leu Arg Ala Cys Lys Lys Gly
50 55

<210> 58
<211> 58
<212> PRT
<213> Artificial Sequence

<220>

<223> DPI.7.2

<400> 58

Gly Pro Ser Trp Cys Leu Thr Pro Ala Asp Arg Gly Leu Cys Val Ala
1 5 10 15
Asn Phe Asn Arg Phe Tyr Tyr Asn Ser Val Ile Gly Lys Cys Arg Pro
20 25 30
Phe Lys Tyr Ser Gly Cys Gly Gly Asn Glu Asn Asn Phe Thr Ser Lys
35 40 45
Gln Glu Cys Leu Arg Ala Cys Lys Lys Gly
50 55

<210> 59
<211> 58
<212> PRT
<213> Artificial Sequence

<220>

<223> DPI.7.3

<400> 59

Gly Pro Ser Trp Cys Leu Thr Pro Ala Asp Arg Gly Leu Cys Val Ala
1 5 10 15

Phe Phe Asn Arg Phe Tyr Tyr Asn Ser Val Ile Gly Lys Cys Arg Pro
20 25 30

Phe Lys Tyr Ser Gly Cys Gly Gly Asn Glu Asn Asn Phe Lys Ser Lys
35 40 45

Gln Glu Cys Leu Arg Ala Cys Lys Lys Gly
50 55

<210> 60
<211> 58
<212> PRT
<213> Artificial Sequence

<220>

<223> DPI.7.4

<400> 60

Gly Pro Ser Trp Cys Leu Thr Pro Ala Val Arg Gly Pro Cys Val Ala
1 5 10 15

Phe Phe Asn Arg Phe Tyr Tyr Asn Ser Val Ile Gly Lys Cys Arg Pro
20 25 30

Phe Lys Tyr Gly Gly Cys Gly Gly Asn Glu Asn Asn Phe Lys Ser Lys
35 40 45

Gln Glu Cys Leu Arg Ala Cys Lys Lys Gly
50 55

<210> 61
<211> 58
<212> PRT
<213> Artificial Sequence

<220>

<223> DPI.7.5

<400> 61

Gly Pro Ser Trp Cys Leu Thr Pro Ala Asp Arg Gly Pro Cys Ile Ala
1 5 10 15

Phe Phe Pro Arg Trp Tyr Tyr Asn Ser Val Ile Gly Lys Cys Gln Thr
20 25 30

Phe Val Tyr Gly Gly Cys Gly Gly Asn Glu Asn Asn Phe Ala Ser Lys
35 40 45

Gln Glu Cys Leu Arg Ala Cys Lys Lys Gly
50 55

<210> 62
<211> 58
<212> PRT

<213> Homo sapiens

<400> 62

Glu Thr Asp Ile Cys Lys Leu Pro Lys Asp Glu Gly Thr Cys Arg Asp
1 5 10 15
Phe Ile Leu Lys Trp Tyr Tyr Asp Pro Asn Thr Lys Ser Cys Ala Arg
20 25 30
Phe Trp Tyr Gly Gly Cys Gly Gly Asn Glu Asn Lys Phe Gly Ser Gln
35 40 45
Lys Glu Cys Glu Lys Val Cys Ala Pro Val
50 55

<210> 63

<211> 58

<212> PRT

<213> Artificial Sequence

<220>

<223> DPI.8.1

<400> 63

Glu Thr Asp Ile Cys Lys Leu Pro Lys Val Arg Gly Pro Cys Ile Ala
1 5 10 15
Phe Phe Pro Arg Trp Tyr Tyr Asp Pro Asn Thr Lys Ser Cys Val Leu
20 25 30
Phe Pro Tyr Gly Gly Cys Gln Gly Asn Gly Asn Lys Phe Gly Ser Gln
35 40 45
Lys Glu Cys Glu Lys Val Cys Ala Pro Val
50 55

<210> 64

<211> 58

<212> PRT

<213> Artificial Sequence

<220>

<223> DPI.8.2

<400> 64

Glu Thr Asp Ile Cys Lys Leu Pro Lys Asp Glu Gly Thr Cys Ile Ala
1 5 10 15
Phe Phe Leu Lys Trp Tyr Tyr Asp Pro Asn Thr Lys Ser Cys Ala Arg
20 25 30
Phe Val Tyr Gly Gly Cys Gly Gly Asn Glu Asn Lys Phe Gly Ser Gln
35 40 45

Lys Glu Cys Glu Lys Val Cys Ala Pro Val
50 55

<210> 65
<211> 58
<212> PRT
<213> Artificial Sequence

<220>

<223> DPI.8.3

<400> 65

Glu Thr Asp Ile Cys Lys Leu Pro Lys Asp Glu Gly Pro Cys Ile Ala
1 5 10 15

Phe Phe Leu Arg Trp Tyr Tyr Asp Pro Asn Thr Lys Ser Cys Ala Arg
20 25 30

Phe Val Tyr Gly Gly Cys Gly Gly Asn Glu Asn Lys Phe Gly Ser Gln
35 40 45

Lys Glu Cys Glu Lys Val Cys Ala Pro Val
50 55

<210> 66
<211> 58
<212> PRT
<213> Homo sapiens

<400> 66

Leu Pro Asn Val Cys Ala Phe Pro Met Glu Lys Gly Pro Cys Gln Thr
1 5 10 15

Tyr Met Thr Arg Trp Phe Phe Asn Phe Glu Thr Gly Glu Cys Glu Leu
20 25 30

Phe Ala Tyr Gly Gly Cys Gly Gly Asn Ser Asn Asn Phe Leu Arg Lys
35 40 45

Glu Lys Cys Glu Lys Phe Cys Lys Phe Thr
50 55

<210> 67
<211> 58
<212> PRT
<213> Artificial Sequence

<220>

<223> DPI.9.1

<400> 67

Leu Pro Asn Val Cys Ala Phe Pro Met Val Arg Gly Pro Cys Ile Ala
1 5 10 15

Phe Phe Pro Arg Trp Phe Phe Asn Phe Glu Thr Gly Glu Cys Val Leu
20 25 30

Phe Val Tyr Gly Gly Cys Gln Gly Asn Gly Asn Asn Phe Leu Arg Lys
35 40 45

Glu Lys Cys Glu Lys Phe Cys Lys Phe Thr
50 55

<210> 68
<211> 58
<212> PRT
<213> Artificial Sequence

<220>

<223> DPI.9.2

<400> 68

Leu Pro Asn Val Cys Ala Phe Pro Met Glu Lys Gly Pro Cys Ile Ala
1 5 10 15

Tyr Phe Thr Arg Trp Phe Phe Asn Phe Glu Thr Gly Glu Cys Glu Leu
20 25 30

Phe Ala Tyr Gly Gly Cys Gly Gly Asn Ser Asn Asn Phe Leu Arg Lys
35 40 45

Glu Lys Cys Glu Lys Phe Cys Lys Phe Thr
50 55

<210> 69
<211> 58
<212> PRT
<213> Artificial Sequence

<220>

<223> DPI.9.3

<400> 69

Leu Pro Asn Val Cys Ala Phe Pro Met Glu Lys Gly Pro Cys Ile Ala
1 5 10 15

Tyr Phe Pro Arg Trp Phe Phe Asn Phe Glu Thr Gly Glu Cys Val Leu
20 25 30

Phe Val Tyr Gly Gly Cys Gly Gly Asn Ser Asn Asn Phe Leu Arg Lys
35 40 45

Glu Lys Cys Glu Lys Phe Cys Lys Phe Thr
50 55

<210> 70
<211> 8157
<212> DNA

<213> Artificial Sequence

<220>

<223> Plasmid pHIL-D2

<400> 70

agatcgcggc cgcgatctaa catccaaaga cgaaagggtg aatgaaacct ttttgccatc	60
cgacatccac aggtccattc tcacacataa gtgccaaacg caacaggagg ggatacacta	120
gcagcagacc gttgcaaacg caggacctcc actcctcttc tcctcaacac ccacttttgc	180
catcgaaaaa ccagcccagt tattgggctt gattggagct cgctcattcc aattccttct	240
attaggctac taacaccatg actttattag cctgtctatc ctggccccc tggcgaggtc	300
atgtttgttt atttccgaat gcaacaagct cgcattaca cccgaacatc actccagatg	360
agggctttct gagtgtggg tcaaatagtt tcatgttccc aaatggcca aaactgacag	420
tttaaacgct gtcttggaac ctaatatgac aaaagcgtga tctcatccaa gatgaactaa	480
gtttggttcg ttgaaatgct aacggccagt tgggtcaaaaa gaaacttcca aaagtcgcca	540
taccgtttgt cttgtttggt attgattgac gaatgctcaa aaataatctc attaatgctt	600
agcgcagtct ctctatcgct tctgaaccog gtggcacctg tgccgaaacg caaatgggga	660
aacaaccgcg tttttgatg attatgcatt gtcctccaca ttgtatgctt ccaagattct	720
ggtgggaata ctgctgatag cctaacgttc atgatcaaaa tttaactgtt ctaaccctta	780
cttgacaggc aatatataaa cagaaggaag ctgccctgtc ttaaaccctt ttttttatca	840
tcattattag cttactttca taattgogac tggttccaat tgacaagctt ttgattttaa	900
cgacttttaa cgacaacttg agaagatcaa aaaacaacta attattcgaa acgaggaatt	960
cgccttagac atgactgttc ctgagttcaa gttgggcatt acgagaagac cggctcttgc	1020
agattctaata caagaggatg tcagaatgcc atttgcctga gagatgcagg cttcattttt	1080
gataactttt tatttgtaac ctatatagta taggattttt tttgtcattt tgtttcttct	1140
cgtacgagct tgctcctgat cagcctatct cgcagctgat gaatatcttg tggtaggggt	1200
ttgggaaaat cattcgagtt tgatgttttt cttggtattt cccactcctc ttcagagtac	1260
agaagattaa gtgagaagtt cgtttggtgca agcttatcga taagctttta tgcggtagtt	1320
tatcacagtt aaattgctaa cgcagtcagg caccgtgtat gaaatctaac aatgcgctca	1380
tcgtcatcct cggcacogtc accctggatg ctgtaggcatt aggcttggtt atgccggtac	1440
tgccgggcct cttgcgggat atcgtccatt ccgacagcat cgccagtcac tatggcgtgc	1500
tgctagcgct atatgcgttg atgcaatttc tatgcgcacc cgttctcgga gcaactgtccg	1560
accgcttttg ccgcgcgcca gtccctgctcg cttecgctact tggagccact atcgactacg	1620

cgatcatggc gaccacaccc gtccgtgtgga tctatcgaat ctaaagttaa gttaaaatct	1680
ctaaataatt aaataagtcc cagtttctcc atacgaacct taacagcatt gcggtgagca	1740
tctagacctt caacagcagc cagatccatc actgcttggc caatatgttt cagtccctca	1800
ggagttacgt cttgtgaagt gatgaacttc tggaagggtg cagtgttaac tccgctgtat	1860
tgacgggcat atccgtacgt tggcaaagtg tggttggtac cggaggagta atctccacaa	1920
ctctctggag agtaggcacc aacaaacaca gatccagcgt gttgtacttg atcaacataa	1980
gaagaagcat tctcgatttg caggatcaag tgttcaggag cgtactgatt ggacatttcc	2040
aaagcctgct cgtaggttgc aaccgatagg gttgtagagt gtgcaatata cttgcgtaca	2100
atttcaaccc ttggcaactg cacagcttgg ttgtgaacag catcttcaat tctggcaagc	2160
tccttgtctg tcatatcgac agccaacaga atcacctggg aatcaatacc atgttcagct	2220
tgagcagaag gtctgaggca acgaaatctg gatcagcgta tttatcagca ataactagaa	2280
cttcagaagg cccagcaggc atgtcaatac tacacagggc tgatgtgtca ttttgaacca	2340
tcattcttggc agcagtaacg aactggtttc ctggaccaa tattttgtca cacttaggaa	2400
cagtttctgt tccgtaagcc atagcagcta ctgcctgggc gcctcctgct agcacgatac	2460
acttagcacc aaccttgtgg gcaacgtaga tgacttctgg ggtaagggtta ccatccttct	2520
taggtggaga tgcaaaaaca atttctttgc aaccagcaac tttggcagga acaccagca	2580
tcagggaagt ggaaggcaga attgcggttc caccaggaat atagaggcca actttctcaa	2640
taggtcttgc aaaacgagag cagactacac cagggaagt ctcaacttgc aacgtctccg	2700
ttagttgagc ttcattggaat ttcttgacgt tatctataga gagatcaatg gctctcttaa	2760
cgttatctgg caattgcata agttcctctg ggaaaggagc ttctaacaca ggtgtcttca	2820
aagcgactcc atcaaaactg gcagttagtt ctaaaagggc tttgtcacca ttttgacgaa	2880
cattgtcgac aattggtttg actaattcca taatctgttc cgttttctgg ataggacgac	2940
gaagggcatc ttcaatttct tgtgaggagg ccttagaaac gtcaattttg cacaattcaa	3000
tacgaccttc agaagggact tcttttaggtt tggattcttc ttaggttgt tccttgggtg	3060
atcctggctt ggcatctcct ttcttcttag tgacctttag ggacttcata tccaggttcc	3120
tctccacctc gtccaacgtc acaccgtact tggcacatct aactaatgca aaataaaata	3180
agtcagcaca ttcccaggct atatcttcct tggatttagc ttctgcaagt tcatcagctt	3240
cctccctaata tttagcgttc aacaaaactt cgtcgtcaaa taaccgtttg gtataagaac	3300
cttctggagc attgctctta cgatcccaca aggtgcttcc atggctctaa gacctttga	3360
ttggccaaaa caggaagtgc gttccaagt acagaaacca acacctgttt gttcaaccac	3420
aaatttcaag cagtctccat cacaatccaa ttcgataccc agcaactttt gagttcgtcc	3480

agatgtagca cctttataacc acaaaccgtg acgacgagat tggtagactc cagtttgtgt	3540
ccttatagcc tccggaatag acttttttga cgagtacacc aggcccaacg agtaattaga	3600
agagtcagcc accaaagtag tgaatagacc atcggggcgg tcagtagtca aagacgccaa	3660
caaaatttca ctgacagga actttttgac atcttcagaa agttcgtatt cagtagtcaa	3720
ttgccgagca tcaataatgg ggattataacc agaagcaaca gtggaagtca catctaccaa	3780
ctttgcggtc tcagaaaaag cataaacagt tctactaccg ccattagtga aacttttcaa	3840
atcgcccagt ggagaagaaa aaggcacagc gatactagca ttagcgggca aggatgcaac	3900
tttatcaacc agggctctat agataaccct agcgctggg atcatccttt ggacaactct	3960
ttctgccaaa tctaggtcca aaatcacttc attgatacca ttatacggat gactcaactt	4020
gcacattaac ttgaagctca gtcgattgag tgaacttgat caggttgtgc agctggtcag	4080
cagcataggg aaacacggct tttcctacca aactcaagga attatcaaac tctgcaacac	4140
ttgcgtatgc aggtagcaag ggaaatgtca tacttgaagt cggacagtga gtgtagtctt	4200
gagaaattct gaagccgtat ttttattatc agtgagtcag tcatcaggag atcctctacg	4260
ccggacgcat cgtggccggc atcacccggc ccacaggtgc ggttgctggc gcctatatcg	4320
ccgacatcac cgatggggaa gatcgggctc gccacttcgg gctcatgagc gcttgtttcg	4380
gcgtgggtat ggtggcaggc cccgtggcg ggggactgtt gggcgccatc tccttgcatg	4440
caccattcct tgcggcggcg gtgctcaacg gcctcaacct actactgggc tgcttcctaa	4500
tgcaggagtc gcataaggga gagcgtcgag tatctatgat tggaagtatg ggaatggtga	4560
taccgcgcat cttcagtgtc ttgaggtctc ctatcagatt atgcccaact aaagcaaccg	4620
gaggaggaga tttcatggta aatttctctg acttttggtc atcagtagac tcgaactgtg	4680
agactatctc ggttatgaca gcagaaatgt ccttcttgga gacagtaaata gaagtccac	4740
caataaagaa atccttggtta tcaggaacaa acttcttggt tcgaactttt tcggtgcctt	4800
gaactataaa atgtagagtg gatatgtcgg gtaggaatgg agcgggcaaa tgcttacctt	4860
ctggaccttc aagaggtatg tagggtttgt agatactgat gccaaacttca gtgacaacgt	4920
tgctatttcg ttcaaaccat tccgaatcca gagaaatcaa agttgtttgt ctactattga	4980
tccaagccag tgcggtcttg aaactgacaa tagtgtgctc gtgttttgag gtcactcttg	5040
tatgaataaa tctagtcttt gatctaaata atcttgacga gccaaaggca taaataccca	5100
aatctaaaac tcttttaaaa cgttaaaagg acaagtatgt ctgcctgtat taaaccccaa	5160
atcagctcgt agtctgatcc tcatcaactt gaggggcact atcttgtttt agagaaattt	5220
gcggagatgc gatatcgaga aaaagggtacg ctgattttaa acgtgaaatt tatctcaaga	5280
tcgcggccgc gatctcgaat aataactgtt atttttcagt gttcccgatc tgcgtctatt	5340

tcacaataacc aacatgagtc agcttatcga tgataagctg tcaaacatga gaattaattc	5400
gatgataagc tgtcaaacat gagaaatctt gaagacgaaa gggcctcgtg atacgcctat	5460
ttttataggt taatgtcatg ataataatgg tttcttagac gtcaggtggc acttttcggg	5520
gaaatgtgcg cggaaccctt atttgtttat ttttctaaat acattcaa atgtatccgc	5580
tcatgagaca ataaccctga taaatgcttc aataatattg aaaaaggaag agtatgagta	5640
ttcaacattt ccgtgtcgcc cttattccct tttttgcggc attttgcctt cctgtttttg	5700
ctcaccaga aacgctggtg aaagtaaaag atgctgaaga tcagttgggt gcacgagtgg	5760
gttacatcga actggatctc aacagcggta agatccttga gagttttcgc cccgaagaac	5820
gttttccaat gatgagcact tttaaagtgc tgctatgtgg cgcggtatta tcccgtgttg	5880
acgccgggca agagcaactc ggtcgccgca tacactattc tcagaatgac ttggttgagt	5940
actcaccagt cacagaaaag catcttacgg atggcatgac agtaagagaa ttatgcagtg	6000
ctgccataac catgagtgat aacactgcgg ccaacttact tctgacaacg atcggaggac	6060
cgaaggagct aaccgctttt ttgcacaaca tgggggatca tgtaactcgc cttgatcgtt	6120
gggaaccgga gctgaatgaa gccataccaa acgacgagcg tgacaccacg atgcctgcag	6180
caatggcaac aacgttgcg aaactattaa ctggcgaact acttactcta gcttcccggc	6240
aacaattaat agactggatg gaggcggata aagttgcagg accacttctg cgctcgcccc	6300
ttccggctgg ctggtttatt gctgataaat ctggagccgg tgagcgtggg tctcgcggta	6360
tcattgcagc actggggcca gatggtaagc cctcccgtat cgtagttatc tacacgacgg	6420
ggagtcaggc aactatggat gaacgaaata gacagatcgc tgagataggt gcctcactga	6480
ttaagcattg gtaactgtca gaccaagttt actcatatat acttttagatt gatttaaatt	6540
gtaaacgtta atattttggt aaaattcgcg tttaaattttt gttaaatcag ctcatttttt	6600
aaccaatagg ccgaaatcgg caaaatccct tataaatcaa agaataagac cgagataggg	6660
ttgagtgttg ttccagtttg gaacaagagt ccactattaa agaacgtgga ctccaacgtc	6720
aaagggcgaa aaaccgtcta tcagggcgat ggcccactac gtgaaccatc accctaataca	6780
agtttttttg ggtcgagggt ccgtaaagca ctaaatcgga accctaaagg gagccccga	6840
tttagagctt gacggggaaa gccggcgaaac gtggcgagaa aggaaggga gaaagcga	6900
ggagcgggcg ctagggcgct ggcaagtgtg gcggtcacgc tgcgcgtaac caccacaccc	6960
gccgcgctta atgcgcgcgt acagggcgcg taaaaggatc taggtgaaga tcctttttga	7020
taatctcatg accaaaatcc cttaacgtga gttttcgctt cactgagcgt cagacccgt	7080
agaaaagatc aaaggatctt cttgagatcc tttttttctg cgcgtaatct gctgcttgca	7140
aacaaaaaaa ccaccgctac cagcgggtgt ttgtttgccg gatcaagagc taccaactct	7200

ttttccgaag gtaactgggt tcagcagagc gcagatacca aatactgtcc ttctagtgtg 7260
gccgtagtta ggccaccact tcaagaactc tgtagcaccg cctacatacc tcgctctgct 7320
aatcctgtta ccagtggctg ctgccagtgg cgataagtcg tgtcttaccc gggttggaactc 7380
aagacgatag ttaccggata aggcgcagcg gtcgggctga acgggggggtt cgtgcacaca 7440
gccagcttg gagcgaacga cctacaccga actgagatac ctacagcgtg agcattgaga 7500
aagcgccacg cttcccgaag ggagaaaggc ggacaggtat ccggt'aagcg gcaggggtcgg 7560
aacaggagag cgcacgaggg agcttccagg gggaaacgcc tggatatcttt atagtcctgt 7620
cgggtttcgc caoctctgac ttgagcgtcg atttttgtga tgctcgtcag gggggcggag 7680
cctatggaaa aacgccagca acgcggcctt ttacgggttc ctggcctttt gctggccttt 7740
tgctcacatg ttctttcctg cgttatcccc tgattctgtg gataaccgta ttaccgcctt 7800
tgagtgagct gataccgctc gccgcagccg aacgaccgag cgcagcgagt cagtgagcga 7860
ggaagcggaa gagcgcctga tgcggtatTT tctccttaag catctgtgcg gtatttcaca 7920
ccgcataatg tgcactctca gtacaatctg ctctgatgcc gcatagttaa gccagtatac 7980
actccgctat cgctacgtga ctgggtcatg gctgcgcccc gacacccgcc aacacccgct 8040
gacgcgccct gacgggcttg tctgctcccg gcatccgctt acagacaagc tgtgaccgtc 8100
tccgggagct gcatgtgtca gaggttttca ccgtcatcac cgaaacgcgc gaggcag 8157

<210> 71

<211> 8584

<212> DNA

<213> Artificial Sequence

<220>

<223> Plasmid pHIL-D2 (MFalphaPrePro::EPI-HNE-3)

<400> 71

agatcgcggc cgcgatctaa catccaaaga cgaaagggtg aatgaaacct ttttgccatc 60
cgacatccac aggtccattc tcacacataa gtgccaaacg caacaggagg ggatacacta 120
gcagcagacc gttgcaaacg caggacctcc actcctcttc tctcaaacac ccacttttgc 180
catcgaaaaa ccagcccagt tattgggctt gattggagct cgctcattcc aattccttct 240
attaggctac taacaccatg actttattag cctgtctatc ctggccccc tggcgaggtc 300
atgtttgttt atttccgaat gcaacaagct ccgcattaca cccgaacatc actccagatg 360
agggttttct gagtgtgggg tcaaatagtt tcatgttccc aaatggccca aaactgacag 420
tttaaacgct gtcttggaac ctaatatgac aaaagcgtga tctcatccaa gatgaactaa 480
gtttggttcg ttgaaatgct aacggccagt tggTcaaaaa gaaacttcca aaagtcgcca 540

taccgtttgt	cttgtttggt	attgattgac	gaatgctcaa	aaataatctc	attaatgctt	600
agcgcagtct	ctctatcgct	tctgaacccg	gtggcacctg	tgccgaaacg	caaatgggga	660
aacaacccgc	tttttgatg	attatgcatt	gtcctccaca	ttgtatgctt	ccaagattct	720
ggtgggaata	ctgctgatag	cctaacgttc	atgatcaaaa	tttaactggt	ctaacccta	780
cttgacaggc	aatatataaa	cagaaggaag	ctgccctgtc	ttaaaccttt	ttttttatca	840
tcattattag	cttactttca	taattgcgac	tggttccaat	tgacaagctt	ttgattttaa	900
cgacttttaa	cgacaacttg	agaagatcaa	aaaacaacta	attattcgaa	acgatgagat	960
tcccatctat	cttcaactgt	gttttgttcg	ctgcttcctc	tgctttggct	gctccagtta	1020
acaccactac	tgaagacgag	actgctcaaa	ttcctgctga	ggctgtcatc	ggttactctg	1080
acttgggaag	tgacttcgac	gtcgtgtttt	tgccattctc	taactctact	aacaacgggt	1140
tgttgttcat	caacactacc	atcgcttcta	tcgctgctaa	ggaggaaggt	gtttccttgg	1200
acaagagagc	tgcttgtaac	ttgccaatcg	tcagagggtc	atgcattgct	ttcttcccaa	1260
gatgggcttt	cgacgctgtt	aagggttaagt	gcgtcttggt	cccatcgggt	ggttgtcaag	1320
gtaacggtaa	caagttctac	tctgagaagg	agtgtagaga	gtactgtggt	gttccatagt	1380
aagaattcgc	cttagacatg	actgttcctc	agttcaagtt	gggcattacg	agaagaccgg	1440
tcttgctaga	ttctaataca	gaggatgtca	gaatgccatt	tgcttgagag	atgcaggctt	1500
catttttgat	acttttttat	ttgtaaccta	tatagtatag	gatttttttt	gtcattttgt	1560
ttcttctcgt	acgagcttgc	tcctgatcag	cctatctcgc	agctgatgaa	tatcttgtgg	1620
taggggtttg	ggaaaatcat	tcgagtttga	tgtttttctt	ggattttccc	actcctcttc	1680
agagtacaga	agattaagtg	agaagttcgt	ttgtgcaagc	ttatcgataa	gctttaatgc	1740
ggtagtttat	cacagttaaa	ttgctaacgc	agtcaggcac	cgtgtatgaa	atctaacaat	1800
gcgtcatcgc	tcctcctcgc	caccgtcacc	ctggatgctg	taggcatagg	cttggttatg	1860
ccggtactgc	cgggcctctt	gcgggatata	gtccattccg	acagcatcgc	cagtcactat	1920
ggcgtgctgc	tagcgtata	tgcgttgatg	caatttctat	gcgcaccctg	tctcggagca	1980
ctgtccgacc	gctttggccg	ccgcccagtc	ctgctcgctt	cgctacttgg	agccactatc	2040
gactacgcga	tcatggcgac	cacaccctgc	ctgtggatct	atcgaatcta	aatgtaagtt	2100
aaaatctcta	aataattaa	taagtcccag	tttctccata	cgaaccttaa	cagcattgcg	2160
gtgagcatct	agaccttcaa	cagcagccag	atccatcact	gcttggccaa	tatgtttcag	2220
tccctcagga	gttacgtctt	gtgaagtgat	gaacttctgg	aaggttgcag	tgtaaactcc	2280
gctgtattga	cgggcatata	cgtacgttgg	caaagtgtgg	ttggtaccgg	aggagtaatc	2340
tccacaactc	tctggagagt	aggcaccaac	aaacacagat	ccagcgtggt	gtacttgatc	2400

aacataagaa	gaagcattct	cgatttgcag	gatcaagtgt	tcaggagcgt	actgattgga	2460
catttccaaa	gcttgctcgt	aggttgcaac	cgataggggt	gtagagtgtg	caatacactt	2520
gcgtacaatt	tcaacccttg	gcaactgcac	agcttggttg	tgaacagcat	cttcaattct	2580
ggcaagctcc	ttgtctgtca	tatcgacagc	caacagaatc	acctgggaat	caataccatg	2640
ttcagcttga	gcagaaggtc	tgaggcaacg	aaatctggat	cagcgtatth	atcagcaata	2700
actagaactt	cagaaggccc	agcaggcatg	tcaatactac	acagggctga	tgtgtcattt	2760
tgaaccatca	tcttggcagc	agtaacgaac	tggtttcctg	gaccaaata	tttgtcacac	2820
ttaggaacag	tttctgttcc	gtaagccata	gcagctactg	cctggggcgcc	tcctgctagc	2880
acgatacact	tagcaccaac	cttgtgggca	acgtagatga	cttctggggg	aagggtacca	2940
tccttcttag	gtggagatgc	aaaaacaatt	tccttgcaac	cagcaactth	ggcaggaaca	3000
cccagcatca	gggaagtggg	aggcagaatt	gcggttccac	caggaatata	gaggccaact	3060
ttctcaatag	gtcttgcaaa	acgagagcag	actacaccag	ggcaagtctc	aacttgcaac	3120
gtctccgtta	gttgagcttc	atggaatttc	ctgacgttat	ctatagagag	atcaatggct	3180
ctcttaacgt	tatctggcaa	ttgcataagt	tcctctggga	aaggagcttc	taacacaggt	3240
gtcttcaaag	cgactccatc	aaacttggca	gttagttcta	aaagggtctt	gtcaccattt	3300
tgacgaacat	tgtcgacaat	tggtttgact	aattccataa	tctgttccgt	tttctggata	3360
ggacgacgaa	gggcatcttc	aatttcttgt	gaggaggcct	tagaaacgtc	aattttgcac	3420
aattcaatac	gaccttcaga	agggaacttc	ttaggtttgg	attcttctth	aggttggtcc	3480
ttggtgtatc	ctggcttggc	atctcctttc	cttctagtga	cctttaggga	cttcatatcc	3540
aggtttctct	ccacctcgtc	caacgtcaca	ccgtacttgg	cacatctaac	taatgcaaaa	3600
taaaataagt	cagcacattc	ccaggctata	tcttccttgg	atttagcttc	tgcaagttca	3660
tcagcttcct	ccctaattth	agcgttcaac	aaaacttcgt	cgtcaaataa	ccgtttggta	3720
taagaacctt	ctggagcatt	gctcttacga	tcccacaagg	tgcttccatg	gctctaagac	3780
cctttgattg	gccaaaacag	gaagtgcggt	ccaagtgaca	gaaaccaaca	cctgtttggt	3840
caaccacaaa	tttcaagcag	tctccatcac	aatccaattc	gataccagc	aacttttgag	3900
ttcgtccaga	tgtagcacct	ttataaccaca	aaccgtgacg	acgagattgg	tagactccag	3960
tttgtgtcct	tatagcctcc	ggaatagact	ttttggacga	gtacaccagg	ccaacgagt	4020
aattagaaga	gtcagccacc	aaagtagtga	atagaccatc	ggggcggtca	gtagtcaaag	4080
acgccaacaa	aatttctactg	acagggaaact	ttttgacatc	ttcagaaagt	tcgtattcag	4140
tagtcaattg	ccgagcatca	ataatgggga	ttataaccaga	agcaacagtg	gaagtacat	4200
ctaccaactt	tgcggtctca	gaaaaagcat	aaacagttct	actaccgcca	ttagtgaaac	4260

ttttcaaadc	gcccagtgga	gaagaaaaag	gcacagcgat	actagcatta	gcgggcaagg	4320
atgcaacttt	atcaaccagg	gtcctataga	taaccctagc	gcctgggata	atccttttga	4380
caactctttc	tgccaaatct	aggtccaaaa	tcacttcatt	gataccatta	tacggatgac	4440
tcaacttgca	cattaacttg	aagctcagtc	gattgagtga	acttgatcag	gttgtgcagc	4500
tggtcagcag	catagggaaa	cacggctttt	cctaccaaac	tcaaggaatt	atcaaactct	4560
gcaacacttg	cgtatgcagg	tagcaaggga	aatgtcatac	ttgaagtcgg	acagtgagtg	4620
tagtcttgag	aaattctgaa	gccgtatttt	tattatcagt	gagtcagtca	tcaggagata	4680
ctctacgccg	gacgcacgt	ggccggcata	accggcgcca	caggtgcggt	tgctggcgcc	4740
tatatcgccg	acatcaccca	tggggaagat	cgggctcgcc	acttcgggct	catgagcgct	4800
tgtttcggcg	tgggtatggt	ggcaggcccc	gtggccgggg	gactgttggg	cgccatctcc	4860
ttgcatgcac	cattccttgc	ggcggcggtg	ctcaacggcc	tcaacctact	actgggctgc	4920
ttcctaatac	aggagtcgca	taaggagag	cgctcagtat	ctatgattgg	aagtatggga	4980
atggtgatac	ccgcattctt	cagtgtcttg	aggtctccta	tcagattatg	cccaactaaa	5040
gcaaccggag	gaggagattt	catggtaaat	ttctctgact	tttggtcata	agtagactcg	5100
aactgtgaga	ctatctcggt	tatgacagca	gaaatgtcct	tcttgagagc	agtaaatgaa	5160
gtcccaccaa	taaagaaatc	cttggttatca	ggaacaaact	tcttgtttcg	aactttttcg	5220
gtgccttgaa	ctataaaatg	tagagtggat	atgtcgggta	ggaatggagc	gggcaaatac	5280
ttaccttctg	gaccttcaag	aggtatgtag	ggtttgtaga	tactgatgcc	aacttcagtg	5340
acaacgttgc	tatttcgttc	aaaccattcc	gaatccagag	aatcaaagt	tgtttgtcta	5400
ctattgatcc	aagccagtgc	ggtcttgaaa	ctgacaatag	tgtgctcgtg	ttttgaggtc	5460
atctttgtat	gaataaatct	agtctttgat	ctaaataatc	ttgacgagcc	aaggcgataa	5520
atacccaaata	ctaaaactct	tttaaaacgt	taaaaggaca	agtatgtctg	cctgtattaa	5580
accccaaata	agctcgtagt	ctgatcctca	tcaacttgag	gggcactata	ttgttttaga	5640
gaaatttgcg	gagatgcgat	atcgagaaaa	aggtacgctg	attttaaacg	tgaaatttat	5700
ctcaagatcg	cggccgcgat	ctcgaataat	aactgttatt	tttcagtgtt	cccgatctgc	5760
gtctatttca	caataccaac	atgagtcagc	ttatcgatga	taagctgtca	aacatgagaa	5820
ttaattcgat	gataagctgt	caaacatgag	aatcttgaa	gacgaaaggg	cctcgtgata	5880
cgcctatttt	tataggttaa	tgtcatgata	ataatggttt	cttagacgtc	aggtggcact	5940
tttcggggaa	atgtgcgcgg	aaccctatt	tgtttatttt	tctaaataca	ttcaaatacg	6000
tatccgctca	tgagacaata	accctgataa	atgcttcaat	aatattgaaa	aaggaagagt	6060
atgagtattc	aacatttccg	tgtcgccctt	attccctttt	ttgcggcatt	ttgccttctt	6120

gtttttgctc	accagaaac	gctggtgaaa	gtaaaagatg	ctgaagatca	gttgggtgca	6180
cgagtgggtt	acatcgaact	ggatctcaac	agcggtaaga	tccttgagag	ttttcgcccc	6240
gaagaacgtt	ttccaatgat	gagcactttt	aaagttctgc	tatgtggcgc	ggtattatcc	6300
cgtgttgacg	cggggcaaga	gcaactcggg	cgccgcatac	actattctca	gaatgacttg	6360
gttgagtact	caccagtcac	agaaaagcat	cttacggatg	gcatgacagt	aagagaatta	6420
tgcagtgctg	ccataaccat	gagtataaac	actgcgcca	acttacttct	gacaacgatc	6480
ggaggaccga	aggagctaac	cgcttttttg	cacaacatgg	gggatcatgt	aactcgcctt	6540
gatcgttggg	aaccggagct	gaatgaagcc	ataccaaacg	acgagcgtga	caccacgatg	6600
cctgcagcaa	tggcaacaac	gttgcgcaaa	ctattaactg	gcgaactact	tactctagct	6660
tcccggcaac	aattaataga	ctggatggag	gcgataaag	ttgcaggacc	acttctgcgc	6720
tggcccttc	cggctggctg	gtttattgct	gataaatctg	gagccggtga	gcgtgggtct	6780
cgcggtatca	ttgcagcact	ggggccagat	ggtaagccct	ccgtatcgt	agttatctac	6840
acgacgggga	gtcaggcaac	tatggatgaa	cgaaatagac	agatcgctga	gatagggtgc	6900
tcactgatta	agcattggta	actgtcagac	caagtttact	catatatact	ttagattgat	6960
ttaaattgta	aacgttaata	ttttgttaaa	attcgcgtta	aatttttggt	aaatcagctc	7020
attttttaac	caataggccg	aaatcggcaa	aatcccttat	aaatcaaaag	aatagaccga	7080
gatagggttg	agtgttgctc	cagtttgga	caagagtcca	ctattaaaga	acgtggactc	7140
caacgtcaaa	gggcgaaaaa	ccgtctatca	gggcgatggc	ccactacgtg	aaccatcacc	7200
ctaatcaagt	ttttggggg	cgaggtgccg	taaagcacta	aatcggaacc	ctaaagggag	7260
ccccgatatt	agagcttgac	ggggaaagcc	ggcgaacgtg	gcgagaaagg	aagggaagaa	7320
agcgaaagga	gcgggcgcta	gggcgctggc	aagtgtagcg	gtcacgctgc	gcgtaaccac	7380
cacaccgcc	gcgttaatg	cgccgctaca	gggcgcgtaa	aaggatctag	gtgaagatcc	7440
tttttgataa	tctcatgacc	aaaatccctt	aacgtgagtt	ttcgttccac	tgagcgtcag	7500
accccgtaga	aaagatcaaa	ggatcttctt	gagatccttt	ttttctgcgc	gtaatctgct	7560
gcttgcaaac	aaaaaaacca	ccgctaccag	cgggtggttg	tttgccggat	caagagctac	7620
caactctttt	tccgaaggta	actggcttca	gcagagcgca	gataccaaat	actgtccttc	7680
tagtgtagcc	gtagttaggc	caccacttca	agaactctgt	agcaccgcct	acatacctcg	7740
ctctgcta	cctgttacca	gtggctgctg	ccagtggcga	taagtcgtgt	cttaccgggt	7800
tggactcaag	acgatagtta	ccggataagg	cgcagcggtc	gggctgaacg	gggggttcgt	7860
gcacacagcc	cagcttggag	cgaacgacct	acaccgaact	gagataccta	cagcgtgagc	7920
attgagaaag	cgccacgctt	cccgaaggga	gaaaggcgga	caggtatccg	gtaagcggca	7980

```

gggtcggaac aggagagcgc acgaggggagc ttccaggggg aaacgcctgg tatctttata 8040
gtcctgtcgg gtttcgccac ctctgacttg agcgtcgatt tttgtgatgc tcgtcagggg 8100
ggcggagcct atggaaaaaac gccagcaacg cggccttttt acggttcctg gccttttct 8160
ggccttttgc tcacatgttc tttcctgctg tatcccctga ttctgtggat aaccgtatta 8220
ccgcctttga gtgagctgat accgctcgcc gcagccgaac gaccgagcgc agcgagtcag 8280
tgagcgagga agcggaagag cgctgatgc ggtattttct ccttacgcat ctgtgcggta 8340
tttcacaccg catatggtgc actctcagta caatctgctc tgatgccgca tagttaagcc 8400
agtatacact ccgctatcgc tacgtgactg ggtcatggct gcgccccgac acccgccaac 8460
acccgctgac gcgccctgac gggcttgtct gctcccgga tccgcttaca gacaagctgt 8520
gaccgtctcc gggagctgca tgtgtcagag gttttcaccg tcatcaccga aacgcgcgag 8580
gcag 8584

```

```

<210> 72
<211> 141
<212> PRT
<213> Artificial Sequence

```

<220>

<223> Plasmid pHIL-D2 (MFalphaPrePro::EPI-HNE-3)

<400> 72

```

Met Arg Phe Pro Ser Ile Phe Thr Ala Val Leu Phe Ala Ala Ser Ser
1          5          10          15

Ala Leu Ala Ala Pro Val Asn Thr Thr Thr Glu Asp Glu Thr Ala Gln
          20          25          30

Ile Pro Ala Glu Ala Val Ile Gly Tyr Ser Asp Leu Glu Gly Asp Phe
          35          40          45

Asp Val Ala Val Leu Pro Phe Ser Asn Ser Thr Asn Asn Gly Leu Leu
          50          55          60

Phe Ile Asn Thr Thr Ile Ala Ser Ile Ala Ala Lys Glu Glu Gly Val
          65          70          75          80

Ser Leu Asp Lys Arg Ala Ala Cys Asn Leu Pro Ile Val Arg Gly Pro
          85          90          95

Cys Ile Ala Phe Phe Pro Arg Trp Ala Phe Asp Ala Val Lys Gly Lys
          100          105          110

Cys Val Leu Phe Pro Tyr Gly Gly Cys Gln Gly Asn Gly Asn Lys Phe
          115          120          125

Tyr Ser Glu Lys Glu Cys Arg Glu Tyr Cys Gly Val Pro
          130          135          140

```

<210> 73
<211> 444
<212> DNA
<213> Artificial Sequence

<220>

<223> BstBI-AatII-EcoRI cassette for expression of Epi-HNE-4

<400> 73
ttcgaacga tgagattccc atctatcttc actgctgttt tggtcgtgc ttcctctgct 60
ttggctgctc cagttaacac cactactgaa gacgagactg ctcaaattcc tgctgaggct 120
gtcatcgggtt actctgactt ggaaggtgac ttcgacgtcg ctgttttgcc attctctaac 180
tctactaaca acggtttgtt gttcatcaac actaccatcg cttctatcgc tgctaaggag 240
gaaggtgttt ccttgacaa gagagaggct tgtaacttgc caatcgtcag aggtccatgc 300
attgctttct tccaagatg ggctttcgac gctgttaagg gtaagtgcgt cttgttccca 360
tacggtggtt gtcaaggtaa cggtacaag ttctactctg agaaggagtg tagagagtac 420
tgtggtgttc catagtaaga attc 444

<210> 74
<211> 141
<212> PRT
<213> Artificial Sequence

<220>

<223> BstBI-AatII-EcoRI cassette for expression of Epi-HNE-4

<400> 74

Met	Arg	Phe	Pro	Ser	Ile	Phe	Thr	Ala	Val	Leu	Phe	Ala	Ala	Ser	Ser
1				5					10					15	
Ala	Leu	Ala	Ala	Pro	Val	Asn	Thr	Thr	Thr	Glu	Asp	Glu	Thr	Ala	Gln
			20					25					30		
Ile	Pro	Ala	Glu	Ala	Val	Ile	Gly	Tyr	Ser	Asp	Leu	Glu	Gly	Asp	Phe
		35					40					45			
Asp	Val	Ala	Val	Leu	Pro	Phe	Ser	Asn	Ser	Thr	Asn	Asn	Gly	Leu	Leu
	50					55					60				
Phe	Ile	Asn	Thr	Thr	Ile	Ala	Ser	Ile	Ala	Ala	Lys	Glu	Glu	Gly	Val
65					70				75					80	
Ser	Leu	Asp	Lys	Arg	Glu	Ala	Cys	Asn	Leu	Pro	Ile	Val	Arg	Gly	Pro
			85						90					95	
Cys	Ile	Ala	Phe	Phe	Pro	Arg	Trp	Ala	Phe	Asp	Ala	Val	Lys	Gly	Lys
		100						105					110		
Cys	Val	Leu	Phe	Pro	Tyr	Gly	Gly	Cys	Gln	Gly	Asn	Gly	Asn	Lys	Phe
		115					120					125			

Tyr Ser Glu Lys Glu Cys Arg Glu Tyr Cys Gly Val Pro
130 135 140

<210> 75
<211> 8590
<212> DNA
<213> Artificial Sequence

<220>

<223> pD2pick (MFalphaPrePro::EPI-NHE-3) circular dsDNA

<400> 75
agatcgcggc cgcgatctaa catccaaaga cgaaagggtg aatgaaacct ttttgccatc 60
cgacatccac aggtccattc tcacacataa gtgccaaacg caacaggagg ggatacacta 120
gcagcagacc gttgcaaacg caggacctcc actcctcttc tcctcaacac ccacttttgc 180
catcgaaaaa ccagcccagt tattgggctt gattggagct cgctcattcc aattccttct 240
attaggctac taacaccatg actttattag cctgtctatc ctggccccc tggcgaggtc 300
atgtttgttt atttccgaat gcaacaagct ccgcattaca cccgaacatc actccagatg 360
agggctttct gagtgtgggg tcaaatagtt tcatgttccc aaatggccca aaactgacag 420
tttaaacgct gtcttggaac ctaatatgac aaaagcgtga tctcatccaa gatgaactaa 480
gtttggttcg ttgaaatgct aacggccagt tgggtcaaaa gaaacttcca aaagtgcgca 540
taccgtttgt cttgtttggt attgattgac gaatgctcaa aaataatctc attaattgctt 600
agcgcagtct ctctatcgct tctgaacccg gtggcacctg tgccgaaacg caaatgggga 660
aacaaccgcg tttttggatg attatgcatt gtcctccaca ttgtatgctt ccaagattct 720
ggtgggaata ctgctgatag cctaacgttc atgatcaaaa tttaactgtt ctaaccctta 780
cttgacaggc aatatataaa cagaaggaag ctgccctgtc ttaaaccctt ttttttatca 840
tcattattag cttactttca taattgcgac tggttccaat tgacaagctt ttgattttaa 900
cgacttttaa cgacaacttg agaagatcaa aaaacaacta attattcgaa acgatgagat 960
tcccatctat cttcactgct gttttgttcg ctgcttcctc tgctttggct gctccagtta 1020
acaccactac tgaagacgag actgctcaaa ttctgctga ggctgtcatc gggtactctg 1080
acttggaagg tgacttcgac gtcgctgttt tgccattctc taactctact aacaacgggt 1140
tgttgttcat caacactacc atcgcttcta tcgctgctaa ggaggaagggt gtttccttgg 1200
acaagagagc tgcttgtaac ttgccaatcg tcagagggtc atgcattgct ttcttcccaa 1260
gatgggcttt cgacgctgtt aagggttaagt gcgtcttggt ccatacgggt ggttgtaag 1320
gtaacggtaa caagttctac tctgagaagg agtgtagaga gtactgtggt gttccatagt 1380
aagaattcgc cttagacatg actgttcctc agttcaagtt gggcattacg agaagaccgg 1440

tcttgctaga ttctaatacaa gaggatgtca gaatgccatt tgccctgagag atgcaggctt	1500
catttttgat acttttttat ttgtaaccta tatagtatag gatttttttt gtcattttgt	1560
ttctttctcgt acgagcttgc tcctgatcag cctatctcgc agctgatgaa tatcttggtg	1620
taggggtttg ggaaaatcat tcgagtttga tgtttttctt ggtatttccc actcctcttc	1680
agagtacaga agattaagtg agaagttcgt ttgtgcaagc ttatcgataa gctttaatgc	1740
ggtagtttat cacagttaaa ttgctaacgc agtcaggcac cgtgtatgaa atctaacaat	1800
gcgctcatcg tcatcctcgg caccgtcacc ctggatgctg taggcatagg cttggttatg	1860
ccggtactgc cgggcctctt gcgggatatc gtccattccg acagcatcgc cagtactat	1920
ggcgtgctgc tagcgctata tgcgttgatg caatttctat gcgcacccgt tctcggagca	1980
ctgtccgacc gctttggccg ccgcccagtc ctgctcgtt cgctacttgg agccactatc	2040
gactacgca tcatggcgac cacaccgtc ctgtggatct atcgaatcta aatgtaagtt	2100
aaaatctcta aataattaaa taagtcccag tttctccata cgaaccttaa cagcattgcg	2160
gtgagcatct agaccttcaa cagcagccag atccatcact gcttggccaa tatgtttcag	2220
tccctcagga gttacgtctt gtgaagtgat gaacttctgg aaggttgcag tgttaactcc	2280
gctgtattga cgggcatatc cgtacgttgg caaagtgtgg ttggtaccgg aggagtaatc	2340
tccacaactc tctggagagt aggcaccaac aaacacagat ccagcgtgtt gtacttgatc	2400
aacataagaa gaagcattct cgatttgcag gatcaagtgt tcaggagcgt actgattgga	2460
catttccaaa gcctgctcgt aggttgcaac cgataggggt gtagagtgtg caatacaatt	2520
gcgtacaatt tcaacccttg gcaactgcac agcttggttg tgaacagcat cttcaattct	2580
ggcaagctcc ttgtctgtca tatcgacagc caacagaatc acctgggaat caataccatg	2640
ttcagcttga gcagaaggtc tgaggcaacg aaatctggat cagcgtattt atcagcaata	2700
actagaactt cagaaggccc agcaggcatg tcaatactac acagggctga tgtgtcattt	2760
tgaaccatca tcttggcagc agtaacgaac tggtttcctg gaccaaata tttgtcacac	2820
ttaggaacag tttctgttcc gtaagccata gcagctactg cctgggcgcc tctgctagc	2880
acgatacact tagcaccaac cttgtgggca acgtagatga cttctgggggt aagggtacca	2940
tccttcttag gtggagatgc aaaaacaatt tctttgcaac cagcaacttt ggcaggaaca	3000
cccagcatca gggaagtgga aggcagaatt gcggttccac caggaatata gaggccaaact	3060
ttctcaatag gtcttgcaaa acgagagcag actacaccag ggcaagtctc aacttgcaac	3120
gtctccgtta gttgagcttc atggaatttc ctgacgttat ctatagagag atcaatggct	3180
ctcttaacgt tatctggcaa ttgcataagt tcctctggga aaggagcttc taacacaggt	3240
gtcttcaaag cgactccatc aaacttggca gttagttcta aaagggttt gtcaccattt	3300

tgacgaacat tgtcgacaat tggtttgact aattccataa tctgttccgt tttctggata	3360
ggacgacgaa gggcatcttc aatttcttgt gaggaggcct tagaaacgtc aattttgcac	3420
aattcaatac gaccttcaga agggacttct ttaggttttg attcttcttt aggttggtcc	3480
ttggtgtatc ctggcttggc atctcctttc cttctagtga ctttaggga cttcatatcc	3540
aggtttctct ccacctcgtc caacgtcaca ccgtacttgg cacatctaac taatgcaaaa	3600
taaaataagt cagcacattc ccaggctata tcttccttgg atttagcttc tgcaagttca	3660
tcagcttcct ccctaatttt agcgttcaac aaaacttcgt cgtcaaataa ccgtttggta	3720
taagaacctt ctggagcatt gctcttacga tcccacaagg tgcttccatg gctctaagac	3780
cctttgattg gccaaaacag gaagtgcgtt ccaagtgaca gaaaccaaca cctgtttggt	3840
caaccacaaa tttcaagcag tctccatcac aatccaattc gataccagc aacttttgag	3900
ttcgtccaga tgtagcacct ttataccaca aaccgtgacg acgagattgg tagactccag	3960
tttgtgtcct tatagcctcc ggaatagact ttttggaaga gtacaccagg cccaacgagt	4020
aattagaaga gtcagccacc aaagtagtga atagaccatc ggggcgggtca gtagtcaaag	4080
acgccaacaa aatttcactg acagggaact ttttgacatc ttcagaaagt tcgtattcag	4140
tagtcaattg ccgagcatca ataatgggga ttataccaga agcaacagtg gaagtcacat	4200
ctaccaactt tgccgtctca gaaaaagcat aaacagttct actaccgcca ttagtgaaac	4260
ttttcaaata gccagtgga gaagaaaaag gcacagcgat actagcatta gcgggcaagg	4320
atgcaacttt atcaaccagg gtcctataga taaccctagc gcctgggatc atcctttgga	4380
caactctttc tgccaaatct aggtccaaaa tcaattcatt gataccatta tacggatgac	4440
tcaacttgca cattaacttg aagctcagtc gattgagtga acttgatcag gttgtgcagc	4500
tggtcagcag catagggaaa cacggctttt cctaccaaac tcaaggaatt atcaaactct	4560
gcaacacttg cgtatgcagg tagcaaggga aatgtcatatc ttgaagtcgg acagtgagtg	4620
tagtcttgag aaattctgaa gccgtatfff tattatcagt gagtcagtca tcaggagatc	4680
ctctacgccg gacgcatcgt ggccggcatc accggcgcca caggtgcggt tgctggcgcc	4740
tatatcgccg acatcaccga tggggaagat cgggctcgcc acttcgggct catgagcgct	4800
tgtttcggcg tgggtatggg ggcaggcccc gtggccgggg gactgttggg cgccatctcc	4860
ttgcatgcac cattccttgc ggccggcggt ctcaacggcc tcaacctact actgggctgc	4920
ttcctaattgc aggagtcgca taaggagag cgtcgagtat ctatgattgg aagtatggga	4980
atggtgatac ccgattctt cagtgtcttg aggtctccta tcagattatg cccaactaaa	5040
gcaaccggag gaggagattt catggtaaat ttctctgact tttggtcac agtagactcg	5100
aactgtgaga ctatctcggt tatgacagca gaaatgtcct tcttgagagac agtaaatgaa	5160

gtccccaccaa	taaagaaatc	cttgttatca	ggaacaaact	tcttgtttcg	cgaacttttt	5220
cggcgccttg	aactataaaa	tgtagagtgg	atatgtcggg	taggaatgga	gcgggcaa	5280
gcttaccttc	tggaccttca	agaggatatg	agggtttgta	gatactgatg	ccaacttcag	5340
tgacaacggt	gctatttcgt	tcaaaccatt	ccgaatccag	agaaatcaaa	gttgtttgtc	5400
tactattgat	ccaagccagt	gcggtcttga	aactgacaat	agtgtgctcg	tgttttgagg	5460
tcatctttgt	atgaataaat	ctagtctttg	atctaaataa	tcttgacgag	ccaaggcgat	5520
aaatacccaa	atctaaaact	cttttaaaac	gttaaaagga	caagtatgtc	tgccctgtatt	5580
aaacccccaaa	tcagctcgta	gtctgatcct	catcaacttg	aggggcacta	tcttgtttta	5640
gagaaatttg	cggagatgcg	atctcgagaa	aaagggtacgc	tgatttttaa	cgtgaaat	5700
atctcaagat	cgcggccgcg	atctcgaaata	ataactgtta	tttttcagt	ttcccgatct	5760
gcgtctat	cacaatacca	acatgagtca	gcttatcgat	gataagctgt	caaacatgag	5820
aattaattcg	atgataagct	gtcaaaca	agaaatcttg	aagacgaaag	ggcctcgtga	5880
tacgcctatt	tttataggtt	aatgtcatga	taataatggt	ttcttagacg	tacgtcaggt	5940
ggcacttttc	ggggaaatgt	gcgcggaacc	cctatttggt	tatttttcta	aatacattca	6000
aatatgtatc	cgctcatgag	acaataaacc	tgataaatgc	ttcaataata	ttgaaaaagg	6060
aagagtatga	gtattcaaca	tttcctgtgc	gcccttattc	ccttttttgc	ggcattttgc	6120
cttctgtttt	ttgctcacc	agaaacgctg	gtgaaagtaa	aagatgctga	agatcagttg	6180
ggtgcacgag	tgggttacat	cgaactggat	ctcaacagcg	gtaagatcct	tgagagt	6240
cgcgccgaag	aacgttttcc	aatgatgagc	acttttaaa	ttctgctatg	tggcgcggt	6300
ttatcccgtg	ttgacgcgg	gcaagagcaa	ctcggtcgcc	gcatacacta	ttctcagaat	6360
gacttggttg	agtactcacc	agtcacagaa	aagcatctta	cggatggcat	gacagtaaga	6420
gaattatgca	gtgctgcat	aaccatgagt	gataaactg	cggccaactt	acttctgaca	6480
acgatcggag	gaccgaagga	gctaaccgct	tttttgcaca	acatggggga	tcatgtaact	6540
cgccttgatc	gttggaacc	ggagctgaat	gaagccatac	caaacgacga	gcgtgacacc	6600
acgatgcctg	cagcaatggc	aacaacgttg	cgcaaactat	taactggcga	actacttact	6660
ctagcttccc	ggcaacaatt	aatagactgg	atggaggcgg	ataaagttgc	aggaccactt	6720
ctgcgctcgg	cccttcggc	tggctggttt	attgctgata	aatctggagc	cggtgagcgt	6780
gggtctcgcg	gtatcattgc	agcactgggg	ccagatggta	agccctcccg	tatcgtagtt	6840
atctacacga	cggggagtca	ggcaactatg	gatgaacgaa	atagacagat	cgctgagata	6900
ggtgcctcac	tgattaagca	ttggtaactg	tcagaccaag	tttactcata	tatactttag	6960
attgatttaa	attgtaaacg	ttaatat	gttaaaattc	gcgttaaatt	tttgttaaat	7020

cagctcattt	tttaaccaat	aggccgaaat	cggcaaaatc	ccttataaat	caaaagaata	7080
gaccgagata	gggttgagt	ttgttccagt	ttggaacaag	agtccactat	taaagaacgt	7140
ggactccaac	gtcaaagggc	gaaaaaccgt	ctatcagggc	gatggcccac	tacgtgaacc	7200
atcaccctaa	tcaagttttt	tggggtcgag	gtgccgtaaa	gcactaaatc	ggaaccctaa	7260
agggagcccc	cgatttagag	cttgacgggg	aaagccggcg	aacgtggcga	gaaaggaagg	7320
gaagaaagcg	aaaggagcgg	gcgctagggc	gctggcaagt	gtagcggcca	cgctgcgcgt	7380
aaccaccaca	cccgccgcgc	ttaatgcgcc	gctacagggc	gcgtaaaagg	atctaggtga	7440
agatcctttt	tgataatctc	atgacaaaaa	tcccttaacg	tgagttttcg	ttccactgag	7500
cgtcagaccc	cgtagaaaag	atcaaaggat	cttcttgaga	tccttttttt	ctgcgcgtaa	7560
tctgctgctt	gcaaacaaaa	aaaccaccgc	taccagcggt	ggtttgtttg	ccggatcaag	7620
agctaccaac	tctttttccg	aaggtaactg	gcttcagcag	agcgagata	ccaaatactg	7680
tccttctagt	gtagccgtag	ttaggccacc	acttcaagaa	ctctgtagca	ccgcctacat	7740
acctcgtctt	gctaatectg	ttaccagtgg	ctgctgccag	tggcgataag	tcgtgtctta	7800
ccgggttga	ctcaagacga	tagttaccgg	ataaggcgca	gcggtcgggc	tgaacggggg	7860
gttcgtgcac	acagcccagc	ttggagcgaa	cgacctacac	cgaactgaga	tacctacagc	7920
gtgagcattg	agaaagcgcc	acgcttcccg	aaggagaaaa	ggcggacagg	tatccggtaa	7980
gcggcagggg	cggaacagga	gagcgcacga	gggagcttcc	agggggaaac	gcctggtatc	8040
tttatagtcc	tgtcggggtt	cgccacctct	gacttgagcg	tcgatttttg	tgatgctcgt	8100
cagggggggc	gagcctatgg	aaaaacgcca	gcaacgcggc	ctttttacgg	ttcctggcct	8160
tttgctggcc	ttttgctcac	atgttctttc	ctgcgttatc	ccctgattct	gtggataaacc	8220
gtattaccgc	ctttgagtga	gctgataacc	ctcgccgcag	ccgaacgacc	gagcgagcgc	8280
agtcagttag	cgaggaagcg	gaagagcgcc	tgatgcggta	ttttctcctt	acgcatctgt	8340
gcggtatttc	acaccgcata	tggtgcactc	tcagtacaat	ctgctctgat	gccgcatagt	8400
taagccagta	tacactccgc	tatcgctacg	tgactgggtc	atggctgcgc	cccgacaccc	8460
gccaacaccc	gctgacgcgc	cctgacgggc	ttgtctgctc	ccggcatccg	cttacagaca	8520
agctgtgacc	gtctccggga	gctgcatgtg	tcagagggtt	tcaccgtcat	caccgaaacg	8580
cgcgaggcag						8590

<210> 76

<211> 141

<212> PRT

<213> Artificial Sequence

<220>

<223> EPI-HNE-3 fusion protein

<400> 76

Met Arg Phe Pro Ser Ile Phe Thr Ala Val Leu Phe Ala Ala Ser Ser
1 5 10 15
Ala Leu Ala Ala Pro Val Asn Thr Thr Thr Glu Asp Glu Thr Ala Gln
20 25 30
Ile Pro Ala Glu Ala Val Ile Gly Tyr Ser Asp Leu Glu Gly Asp Phe
35 40 45
Asp Val Ala Val Leu Pro Phe Ser Asn Ser Thr Asn Asn Gly Leu Leu
50 55 60
Phe Ile Asn Thr Thr Ile Ala Ser Ile Ala Ala Lys Glu Glu Gly Val
65 70 75 80
Ser Leu Asp Lys Arg Ala Ala Cys Asn Leu Pro Ile Val Arg Gly Pro
85 90 95
Cys Ile Ala Phe Phe Pro Arg Trp Ala Phe Asp Ala Val Lys Gly Lys
100 105 110
Cys Val Leu Phe Pro Tyr Gly Gly Cys Gln Gly Asn Gly Asn Lys Phe
115 120 125
Tyr Ser Glu Lys Glu Cys Arg Glu Tyr Cys Gly Val Pro
130 135 140

<210> 77

<211> 147

<212> PRT

<213> Homo sapiens

<400> 77

Ala Val Leu Pro Gln Glu Glu Glu Gly Ser Gly Gly Gly Gln Leu Val
1 5 10 15
Thr Glu Val Thr Lys Lys Glu Asp Ser Cys Gln Leu Gly Tyr Ser Ala
20 25 30
Gly Pro Cys Met Gly Met Thr Ser Arg Tyr Phe Tyr Asn Gly Thr Ser
35 40 45
Met Ala Cys Glu Thr Phe Gln Tyr Gly Gly Cys Met Gly Asn Gly Asn
50 55 60
Asn Phe Val Thr Glu Lys Glu Cys Leu Gln Thr Cys Arg Thr Val Ala
65 70 75 80
Ala Cys Asn Leu Pro Ile Val Arg Gly Pro Cys Arg Ala Phe Ile Gln
85 90 95
Leu Trp Ala Phe Asp Ala Val Lys Gly Lys Cys Val Leu Phe Pro Tyr
100 105 110

Gly Gly Cys Gln Gly Asn Gly Asn Lys Phe Tyr Ser Glu Lys Glu Cys
115 120 125

Arg Glu Tyr Cys Gly Val Pro Gly Asp Gly Asp Glu Glu Leu Leu Arg
130 135 140

Phe Ser Asn
145

<210> 78
<211> 249
<212> DNA
<213> Artificial Sequence

<220>

<223> M13_III_signal::Human_LACI-D2::mature_M13_III

<400> 78
atgaagaagc ttctcttcgc cattcctctg gtggtacctt tctattccgg cgccaagcct 60
gacttctgct tctctgagga ggatcccggg atttgccgcg gttatattac gcgttatttc 120
tataataacc agactaagca atgtgagcgg ttcaagtatg gtggttgcct aggtaatatg 180
aacaacttcg agactctaga agagtgtgaag aacatatgtg aggatggtgg tgctgagact 240
gttgagtct 249

<210> 79
<211> 83
<212> PRT
<213> Artificial Sequence

<220>

<223> LACI-D2 fusion protein

<400> 79

Met Lys Lys Leu Leu Phe Ala Ile Pro Leu Val Val Pro Phe Tyr Ser
1 5 10 15

Gly Ala Lys Pro Asp Phe Cys Phe Leu Glu Glu Asp Pro Gly Ile Cys
20 25 30

Arg Gly Tyr Ile Thr Arg Tyr Phe Tyr Asn Asn Gln Thr Lys Gln Cys
35 40 45

Glu Arg Phe Lys Tyr Gly Gly Cys Leu Gly Asn Met Asn Asn Phe Glu
50 55 60

Thr Leu Glu Glu Cys Lys Asn Ile Cys Glu Asp Gly Gly Ala Glu Thr
65 70 75 80

Val Glu Ser

<210> 80
 <211> 189
 <212> DNA
 <213> Artificial Sequence

<220>

<223> laci-d1 with cloning sites

<400> 80
 gcgggccgaga tgcattcctt ctgcgctttc aaagctgatg acgggccgtg taaagctatc 60
 atgaaacggt tcttcttcaa cattttcacg cgtcagtgcg aggaattcat ttacgggtggt 120
 tgtgaaggta accagaaccg gttcgaatct ctagaggaat gtaagaagat gtgcactcgt 180
 gacggcgcc 189

<210> 81
 <211> 63
 <212> PRT
 <213> Artificial Sequence

<220>

<223> laci-d1 with linkers

<400> 81
 Ala Ala Glu Met His Ser Phe Cys Ala Phe Lys Ala Asp Asp Gly Pro
 1 5 10 15
 Cys Lys Ala Ile Met Lys Arg Phe Phe Phe Asn Ile Phe Thr Arg Gln
 20 25 30
 Cys Glu Glu Phe Ile Tyr Gly Gly Cys Glu Gly Asn Gln Asn Arg Phe
 35 40 45
 Glu Ser Leu Glu Glu Cys Lys Lys Met Cys Thr Arg Asp Gly Ala
 50 55 60

<210> 82
 <211> 189
 <212> DNA
 <213> Artificial sequence

<220>

<223> LACI-D1 hNE library

<220>
 <221> misc_feature
 <222> (37)..(37)
 <223> n is a, c, g, or t

<220>
 <221> misc_feature
 <222> (38)..(38)
 <223> n is a or g

<220>
<221> misc_feature
<222> (40)..(40)
<223> n is a or g

<220>
<221> misc_feature
<222> (41)..(41)
<223> n is a, c or g

<220>
<221> misc_feature
<222> (42)..(42)
<223> n is c or g

<220>
<221> misc_feature
<222> (47)..(47)
<223> n is a, c, g, or t

<220>
<221> misc_feature
<222> (52)..(52)
<223> n is a or g

<220>
<221> misc_feature
<222> (56)..(56)
<223> n is c or g

<220>
<221> misc_feature
<222> (58)..(58)
<223> n is a, c, g, or t

<220>
<221> misc_feature
<222> (64)..(64)
<223> n is a or c

<220>
<221> misc_feature
<222> (65)..(65)
<223> n is a, c, g, or t

<220>
<221> misc_feature
<222> (66)..(66)
<223> n is c or g

<220>
<221> misc_feature
<222> (71)..(71)
<223> n is a, g or t

<220>
<221> misc_feature
<222> (72)..(72)
<223> n is c or g

<220>

<221> misc_feature
<222> (100)..(100)
<223> n is c or g

<220>
<221> misc_feature
<222> (101)..(101)
<223> n is a or t

<220>
<221> misc_feature
<222> (103)..(103)
<223> n is a, c or g

<220>
<221> misc_feature
<222> (104)..(104)
<223> n is a, c or t

<220>
<221> misc_feature
<222> (109)..(109)
<223> n is a, c or g

<220>
<221> misc_feature
<222> (110)..(110)
<223> n is a, c or t

<220>
<221> misc_feature
<222> (124)..(124)
<223> n is a, c or g

<220>
<221> misc_feature
<222> (125)..(125)
<223> n is a, c or t

<220>
<221> misc_feature
<222> (128)..(128)
<223> n is c or g

<220>
<221> misc_feature
<222> (133)..(133)
<223> n is c or g

<220>
<221> misc_feature
<222> (134)..(134)
<223> n is a or g

<400> 82
gcggccgaga tgcattcctt ctgcgctttc aaagctnntn nnggtcnttg tnttgntntc 60
ttcnnncggt nnttcttcaa cattttcacg cgtcagtgcg ngnnattcnn atacggtggt 120
tgtngngnta acnngaaccg gttcgaatct ctagaggaat gtaagaagat gtgcactcgt 180

gacggcgcc

189

<210> 83
<211> 63
<212> PRT
<213> Artificial Sequence

<220>

<223> LACI-D1 hNE library

<220>
<221> misc_feature
<222> (13)..(13)
<223> Xaa is Cys, Arg, Ser, Gly, Tyr, His, Asp or Asn

<220>
<221> misc_feature
<222> (14)..(14)
<223> Xaa is Thr, Asn, Lys, Arg, Ser, Ala, Glu, Gly or Asp

<220>
<221> misc_feature
<222> (16)..(16)
<223> Xaa is His, Arg, Pro or Leu

<220>
<221> misc_feature
<222> (18)..(18)
<223> Xaa is Val or Ile

<220>
<221> misc_feature
<222> (19)..(19)
<223> Xaa is Ala or Gly

<220>
<221> misc_feature
<222> (20)..(20)
<223> Xaa is Phe, Leu, Ile or Val

<220>
<221> misc_feature
<222> (22)..(22)
<223> Xaa is Ser, Thr, Asn, Ile, Met, Gln, His, Leu, Pro, Lys or Arg

<220>
<221> misc_feature
<222> (24)..(24)
<223> Xaa is Cys, Tyr, Trp, Phe or Leu

<220>
<221> misc_feature
<222> (34)..(34)
<223> Xaa is Leu, Gln, Glu or Val

<220>
<221> misc_feature
<222> (35)..(35)
<223> Xaa is Gln, Leu, Pro, Thr, Lys, Val, Ile, Glu or Ala

<220>
<221> misc_feature
<222> (37)..(37)
<223> Xaa is Gln, Leu, Pro, Thr, Lys, Val, Glu, Ile or Ala

<220>
<221> misc_feature
<222> (42)..(42)
<223> Xaa is Gln, Leu, Pro, Thr, Lys, Val, Met, Glu or Ala

<220>
<221> misc_feature
<222> (43)..(43)
<223> Xaa is Gly or Ala

<220>
<221> misc_feature
<222> (45)..(45)
<223> Xaa is Glu, Gly, Gln or Arg

<400> 83

Ala	Ala	Glu	Met	His	Ser	Phe	Cys	Ala	Phe	Lys	Ala	Xaa	Xaa	Gly	Xaa
1				5					10					15	

Cys	Xaa	Xaa	Xaa	Phe	Xaa	Arg	Xaa	Phe	Phe	Asn	Ile	Phe	Thr	Arg	Gln
			20				25						30		

Cys	Xaa	Xaa	Phe	Xaa	Tyr	Gly	Gly	Cys	Xaa	Xaa	Asn	Xaa	Asn	Arg	Phe
			35			40						45			

Glu	Ser	Leu	Glu	Glu	Cys	Lys	Lys	Met	Cys	Thr	Arg	Asp	Gly	Ala
	50					55					60			

<210> 84
<211> 201
<212> DNA
<213> Artificial sequence

<220>

<223> LACI-D2 hNE library

<220>
<221> misc_feature
<222> (34)..(34)
<223> n is a, c, g, or t

<220>
<221> misc_feature
<222> (35)..(35)
<223> n is a or g

<220>
<221> misc_feature
<222> (37)..(38)
<223> n is a, c or g

<220>
<221> misc_feature

<222> (39)..(39)
<223> n is c or g

<220>
<221> misc_feature
<222> (43)..(43)
<223> n is a or c

<220>
<221> misc_feature
<222> (44)..(44)
<223> n is a, c, g, or t

<220>
<221> misc_feature
<222> (49)..(49)
<223> n is a or g

<220>
<221> misc_feature
<222> (53)..(53)
<223> n is c or g

<220>
<221> misc_feature
<222> (55)..(55)
<223> n is a, c, g, or t

<220>
<221> misc_feature
<222> (56)..(56)
<223> n is a or t

<220>
<221> misc_feature
<222> (61)..(61)
<223> n is a or c

<220>
<221> misc_feature
<222> (62)..(62)
<223> n is a, c, g, or t

<220>
<221> misc_feature
<222> (63)..(63)
<223> n is c or g

<220>
<221> misc_feature
<222> (68)..(68)
<223> n is a, g or t

<220>
<221> misc_feature
<222> (69)..(69)
<223> n is c or g

<220>
<221> misc_feature
<222> (97)..(97)

<223> n is c or g

<220>

<221> misc_feature

<222> (98)..(98)

<223> n is a or t

<220>

<221> misc_feature

<222> (100)..(100)

<223> n is a, c or g

<220>

<221> misc_feature

<222> (101)..(101)

<223> n is a, c, g, or t

<220>

<221> misc_feature

<222> (106)..(106)

<223> n is a, c, or g

<220>

<221> misc_feature

<222> (107)..(107)

<223> n is a, c, or t

<220>

<221> misc_feature

<222> (121)..(121)

<223> n is a, c, or g

<220>

<221> misc_feature

<222> (122)..(122)

<223> n is a, c, or t

<220>

<221> misc_feature

<222> (125)..(125)

<223> n is c or g

<220>

<221> misc_feature

<222> (130)..(130)

<223> n is a, c or g

<220>

<221> misc_feature

<222> (131)..(131)

<223> n is c, g or t

<400> 84

ggcgccaagc ctgacttctg cttcctcgag gagnntnnng ggnnttgcnt tgntnntttt 60

nnncgttnnt tctataataa ccaggctaag caatgtnnngn nattcnata tgggtggtgc 120

nnngntaatn ngaacaactt cgagactcta gaagagtgtg agaacatatg tgaggatggt 180

ggtgctgaga ctgttgagtc t 201

<210> 85
<211> 67
<212> PRT
<213> Artificial Sequence

<220>

<223> LACI-D2 hNE library

<220>

<221> misc_feature

<222> (12)..(12)

<223> Xaa is Cys, Arg, Ser, Gly, Tyr, His, Asp or Asn

<220>

<221> misc_feature

<222> (13)..(13)

<223> Xaa is Pro, His, Thr, Asn, Lys, Arg, Ser, Ala, Glu, Gly, Asp or Gln

<220>

<221> misc_feature

<222> (15)..(15)

<223> Xaa is His, Arg, Pro, Leu, Asn, Ser, Ile or Thr

<220>

<221> misc_feature

<222> (17)..(17)

<223> Xaa is Val or Ile

<220>

<221> misc_feature

<222> (18)..(18)

<223> Xaa is Gly or Ala

<220>

<221> misc_feature

<222> (19)..(19)

<223> Xaa is Phe, Leu, Ile, Val, Tyr, His, Asn or Asp

<220>

<221> misc_feature

<222> (21)..(21)

<223> Xaa is Ile, Asn, Gln, Met, Leu, His, Lys, Pro, Thr or Arg

<220>

<221> misc_feature

<222> (23)..(23)

<223> Xaa is Cys, Phe, Leu, Tyr or Trp

<220>

<221> misc_feature

<222> (33)..(33)

<223> Xaa is Leu, Gln, Glu or Val

<220>

<221> misc_feature

<222> (34)..(34)

<223> Xaa is Gln, Gly, Leu, Pro, Thr, Lys, Val, Ile, Glu, Ala or Arg

<220>
<221> misc_feature
<222> (36)..(36)
<223> Xaa is Gln, Leu, Pro, Thr, Val, Glu, Ile, Ala or Lys

<220>
<221> misc_feature
<222> (41)..(41)
<223> Xaa is Gln, Pro, Thr, Lys, Val, Met, Glu, Ala or Leu

<220>
<221> misc_feature
<222> (42)..(42)
<223> Xaa is Gly or Ala

<220>
<221> misc_feature
<222> (44)..(44)
<223> Xaa is Arg, Gly, Lys, Glu, Leu, Gln, Met or Val

<400> 85

Gly Ala Lys Pro Asp Phe Cys Phe Leu Glu Glu Xaa Xaa Gly Xaa Cys
1 5 10 15

Xaa Xaa Xaa Phe Xaa Arg Xaa Phe Tyr Asn Asn Gln Ala Lys Gln Cys
20 25 30

Xaa Xaa Phe Xaa Tyr Gly Gly Cys Xaa Xaa Asn Xaa Asn Asn Phe Glu
35 40 45

Thr Leu Glu Glu Cys Lys Asn Ile Cys Glu Asp Gly Gly Ala Glu Thr
50 55 60

Val Glu Ser
65

<210> 86
<211> 51
<212> PRT
<213> Artificial Sequence

<220>

<223> definition of aprotonin-like Kunitz domain (p. 11)

<220>
<221> misc_feature
<222> (2)..(7)
<223> Xaa is any amino acid

<220>
<221> misc_feature
<222> (9)..(9)
<223> Xaa is any amino acid

<220>
<221> misc_feature

<222> (11)..(18)
<223> Xaa is any amino acid

<220>
<221> misc_feature
<222> (19)..(19)
<223> Xaa is any Tyr or Phe

<220>
<221> misc_feature
<222> (20)..(25)
<223> Xaa is any amino acid

<220>
<221> misc_feature
<222> (27)..(28)
<223> Xaa is any amino acid

<220>
<221> misc_feature
<222> (30)..(30)
<223> Xaa is any amino acid
<220>
<221> misc_feature
<222> (31)..(31)
<223> Xaa is Tyr, Trp or Phe

<220>
<221> misc_feature
<222> (32)..(32)
<223> Xaa is any amino acid

<220>
<221> misc_feature
<222> (35)..(38)
<223> Xaa is any amino acid

<220>
<221> misc_feature
<222> (39)..(39)
<223> Xaa is Asn or Gly

<220>
<221> misc_feature
<222> (40)..(40)
<223> Xaa is any amino acid

<220>
<221> misc_feature
<222> (41)..(41)
<223> Xaa is Phe or Tyr

<220>
<221> misc_feature
<222> (42)..(46)
<223> Xaa is any amino acid

<220>
<221> misc_feature
<222> (48)..(50)
<223> Xaa is any amino acid

<400> 86

Cys Xaa Xaa Xaa Xaa Xaa Xaa Gly Xaa Cys Xaa Xaa Xaa Xaa Xaa Xaa
1 5 10 15
Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Cys Xaa Xaa Phe Xaa Xaa Xaa
20 25 30
Gly Cys Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Cys Xaa
35 40 45
Xaa Xaa Cys
50

<210> 87

<211> 58

<212> PRT

<213> Bos Taurus

<400> 87

Arg Pro Asp Phe Cys Leu Glu Pro Pro Tyr Thr Gly Pro Cys Lys Ala
1 5 10 15
Arg Ile Ile Arg Tyr Phe Tyr Asn Ala Lys Ala Gly Leu Cys Gln Thr
20 25 30
Phe Val Tyr Gly Gly Cys Arg Ala Lys Arg Asn Asn Phe Lys Ser Ala
35 40 45
Glu Asp Cys Met Arg Thr Cys Gly Gly Ala
50 55

<210> 88

<211> 58

<212> PRT

<213> Artificial Sequence

<220>

<223> Engineered B-PTI from MARK87

<400> 88

Arg Pro Asp Phe Cys Leu Glu Pro Pro Tyr Thr Gly Pro Thr Lys Ala
1 5 10 15
Arg Ile Ile Arg Tyr Phe Tyr Asn Ala Lys Ala Gly Leu Cys Gln Thr
20 25 30
Phe Val Tyr Gly Gly Thr Arg Ala Lys Arg Asn Asn Phe Lys Ser Ala
35 40 45
Glu Asp Cys Met Arg Thr Cys Gly Gly Ala
50 55

<210> 89
<211> 58
<212> PRT
<213> Artificial Sequence

<220>

<223> Engineered B-PTI from MARK87

<400> 89

Arg Pro Asp Phe Cys Leu Glu Pro Pro Tyr Thr Gly Pro Ala Lys Ala
1 5 10 15
Arg Ile Ile Arg Tyr Phe Tyr Asn Ala Lys Ala Gly Leu Cys Gln Thr
20 25 30
Phe Val Tyr Gly Gly Ala Arg Ala Lys Arg Asn Asn Phe Lys Ser Ala
35 40 45
Glu Asp Cys Met Arg Thr Cys Gly Gly Ala
50 55

<210> 90
<211> 67
<212> PRT
<213> Bos taurus (Bovine Colostrum)

<400> 90

Phe Gln Thr Pro Pro Asp Leu Cys Gln Leu Pro Gln Ala Arg Gly Pro
1 5 10 15
Cys Lys Ala Ala Leu Leu Arg Tyr Phe Tyr Asn Ser Thr Ser Asn Ala
20 25 30
Cys Glu Pro Phe Thr Tyr Gly Gly Cys Gln Gly Asn Asn Asn Asn Phe
35 40 45
Glu Thr Thr Glu Met Cys Leu Arg Ile Cys Glu Pro Pro Gln Gln Thr
50 55 60
Asp Lys Ser
65

<210> 91
<211> 60
<212> PRT
<213> Bos Taurus (Bovine serum)

<400> 91

Thr Glu Arg Pro Asp Phe Cys Leu Glu Pro Pro Tyr Thr Gly Pro Cys
1 5 10 15
Lys Ala Ala Met Ile Arg Tyr Phe Tyr Asn Ala Lys Ala Gly Phe Cys
20 25 30

Glu Thr Phe Val Tyr Gly Gly Cys Arg Ala Lys Ser Asn Asn Phe Lys
35 40 45
Ser Ala Glu Asp Cys Met Arg Thr Cys Gly Gly Ala
50 55 60

<210> 92
<211> 58
<212> PRT
<213> Artificial Sequence

<220>

<223> Semisynthetic BPTI, TSCH87

<400> 92

Arg Pro Asp Phe Cys Leu Glu Pro Pro Tyr Thr Gly Pro Cys Val Ala
1 5 10 15
Arg Ile Ile Arg Tyr Phe Tyr Asn Ala Lys Ala Gly Leu Cys Gln Thr
20 25 30

Phe Val Tyr Gly Gly Cys Arg Ala Lys Arg Asn Asn Phe Lys Ser Ala
35 40 45
Glu Asp Cys Met Arg Thr Cys Gly Gly Ala
50 55

<210> 93
<211> 58
<212> PRT
<213> Artificial Sequence

<220>

<223> Semisynthetic BPTI, TSCH87

<400> 93

Arg Pro Asp Phe Cys Leu Glu Pro Pro Tyr Thr Gly Pro Cys Gly Ala
1 5 10 15
Arg Ile Ile Arg Tyr Phe Tyr Asn Ala Lys Ala Gly Leu Cys Gln Thr
20 25 30

Phe Val Tyr Gly Gly Cys Arg Ala Lys Arg Asn Asn Phe Lys Ser Ala
35 40 45
Glu Asp Cys Met Arg Thr Cys Gly Gly Ala
50 55

<210> 94
<211> 58
<212> PRT
<213> Artificial Sequence

<220>

<223> Semisynthetic BPTI, TSCH87

<400> 94

Arg Pro Asp Phe Cys Leu Glu Pro Pro Tyr Thr Gly Pro Cys Ala Ala
1 5 10 15
Arg Ile Ile Arg Tyr Phe Tyr Asn Ala Lys Ala Gly Leu Cys Gln Thr
20 25 30
Phe Val Tyr Gly Gly Cys Arg Ala Lys Arg Asn Asn Phe Lys Ser Ala
35 40 45
Glu Asp Cys Met Arg Thr Cys Gly Gly Ala
50 55

<210> 95

<211> 58

<212> PRT

<213> Artificial Sequence

<220>

<223> Semisynthetic BPTI, TSCH87

<400> 95

Arg Pro Asp Phe Cys Leu Glu Pro Pro Tyr Thr Gly Pro Cys Leu Ala
1 5 10 15
Arg Ile Ile Arg Tyr Phe Tyr Asn Ala Lys Ala Gly Leu Cys Gln Thr
20 25 30
Phe Val Tyr Gly Gly Cys Arg Ala Lys Arg Asn Asn Phe Lys Ser Ala
35 40 45
Glu Asp Cys Met Arg Thr Cys Gly Gly Ala
50 55

<210> 96

<211> 58

<212> PRT

<213> Artificial Sequence

<220>

<223> Semisynthetic BPTI, TSCH87

<400> 96

Arg Pro Asp Phe Cys Leu Glu Pro Pro Tyr Thr Gly Pro Cys Ile Ala
1 5 10 15
Arg Ile Ile Arg Tyr Phe Tyr Asn Ala Lys Ala Gly Leu Cys Gln Thr
20 25 30

Phe Val Tyr Gly Gly Cys Arg Ala Lys Arg Asn Asn Phe Lys Ser Ala
35 40 45

Glu Asp Cys Met Arg Thr Cys Gly Gly Ala
50 55

<210> 97
<211> 58
<212> PRT
<213> Artificial Sequence

<220>

<223> Engineered BPTI, AUER87

<400> 97

Arg Pro Asp Phe Cys Leu Glu Pro Pro Tyr Thr Gly Pro Cys Lys Ala
1 5 10 15

Arg Ile Ile Arg Tyr Phe Tyr Asn Ala Lys Ala Gly Leu Cys Gln Thr
20 25 30

Phe Val Tyr Gly Gly Cys Arg Ala Lys Arg Asn Asn Phe Lys Ser Ala
35 40 45

Glu Asp Cys Glu Arg Thr Cys Gly Gly Ala
50 55

<210> 98
<211> 60
<212> PRT
<213> Dendroaspis polylepis polylepis (Black mamba venom I)

<400> 98

Gln Pro Leu Arg Lys Leu Cys Ile Leu His Arg Asn Pro Gly Arg Cys
1 5 10 15

Tyr Gln Lys Ile Pro Ala Phe Tyr Tyr Asn Gln Lys Lys Lys Gln Cys
20 25 30

Glu Gly Phe Thr Trp Ser Gly Cys Gly Gly Asn Ser Asn Arg Phe Lys
35 40 45

Thr Ile Glu Glu Cys Arg Arg Thr Cys Ile Arg Lys
50 55 60

<210> 99
<211> 57
<212> PRT
<213> Dendroaspis polylepis polylepis (Black mamba venom K)

<400> 99

Ala Ala Lys Tyr Cys Lys Leu Pro Leu Arg Ile Gly Pro Cys Lys Arg
1 5 10 15

Lys Ile Pro Ser Phe Tyr Tyr Lys Trp Lys Ala Lys Gln Cys Leu Pro
20 25 30

Phe Asp Tyr Ser Gly Cys Gly Gly Asn Ala Asn Arg Phe Lys Thr Ile
35 40 45

Glu Glu Cys Arg Arg Thr Cys Val Gly
50 55

<210> 100
<211> 57
<212> PRT
<213> Hemachatus hemachates

<400> 100

Arg Pro Asp Phe Cys Glu Leu Pro Ala Glu Thr Gly Leu Cys Lys Ala
1 5 10 15

Tyr Ile Arg Ser Phe His Tyr Asn Leu Ala Ala Gln Gln Cys Leu Gln
20 25 30

Phe Ile Tyr Gly Gly Cys Gly Gly Asn Ala Asn Arg Phe Lys Thr Ile
35 40 45

Asp Glu Cys Arg Arg Thr Cys Val Gly
50 55

<210> 101
<211> 57
<212> PRT
<213> Naja nivea

<400> 101

Arg Pro Arg Phe Cys Glu Leu Pro Ala Glu Thr Gly Leu Cys Lys Ala
1 5 10 15

Arg Ile Arg Ser Phe His Tyr Asn Arg Ala Ala Gln Gln Cys Leu Glu
20 25 30

Phe Ile Tyr Gly Gly Cys Gly Gly Asn Ala Asn Arg Phe Lys Thr Ile
35 40 45

Asp Glu Cys His Arg Thr Cys Val Gly
50 55

<210> 102
<211> 60
<212> PRT
<213> Vipera russelli

<400> 102

His Asp Arg Pro Thr Phe Cys Asn Leu Pro Pro Glu Ser Gly Arg Cys
1 5 10 15

Arg Gly His Ile Arg Arg Ile Tyr Tyr Asn Leu Glu Ser Asn Lys Cys
20 25 30
Lys Val Phe Phe Tyr Gly Gly Cys Gly Gly Asn Ala Asn Asn Phe Glu
35 40 45
Thr Arg Asp Glu Cys Arg Glu Thr Cys Gly Gly Lys
50 55 60

<210> 103
<211> 64
<212> PRT
<213> Caretta sp. (Red sea turtle egg white)

<220>
<221> misc_feature
<222> (1)..(1)
<223> Xaa is Glu or Gln

<400> 103

Xaa Gly Asp Lys Arg Asp Ile Cys Arg Leu Pro Pro Glu Gln Gly Pro
1 5 10 15
Cys Lys Gly Arg Leu Pro Arg Tyr Phe Tyr Asn Pro Ala Ser Arg Met
20 25 30
Cys Glu Ser Phe Ile Tyr Gly Gly Cys Lys Gly Asn Lys Asn Asn Phe
35 40 45
Lys Thr Lys Ala Glu Cys Val Arg Ala Cys Arg Pro Pro Glu Arg Pro
50 55 60

<210> 104
<211> 58
<212> PRT
<213> Helix pomania

<220>
<221> misc_feature
<222> (1)..(1)
<223> Xaa is Glu or Gln

<400> 104

Xaa Gly Arg Pro Ser Phe Cys Asn Leu Pro Ala Glu Thr Gly Pro Cys
1 5 10 15
Lys Ala Ser Ile Arg Gln Tyr Tyr Tyr Asn Ser Lys Ser Gly Gly Cys
20 25 30
Gln Gln Phe Ile Tyr Gly Gly Cys Arg Gly Asn Gln Asn Arg Phe Asp
35 40 45
Thr Thr Gln Gln Cys Gln Gly Val Cys Val
50 55

<210> 105
<211> 57
<212> PRT
<213> Dendroaspis angusticeps (Eastern green mamba C13 S1 C3 toxin)

<400> 105

Ala	Ala	Lys	Tyr	Cys	Lys	Leu	Pro	Val	Arg	Tyr	Gly	Pro	Cys	Lys	Lys
1				5					10					15	
Lys	Phe	Pro	Ser	Phe	Tyr	Tyr	Asn	Trp	Lys	Ala	Lys	Gln	Cys	Leu	Pro
			20					25					30		
Phe	Asn	Tyr	Ser	Gly	Cys	Gly	Gly	Asn	Ala	Asn	Arg	Phe	Lys	Thr	Ile
		35					40					45			
Glu	Glu	Cys	Arg	Arg	Thr	Cys	Val	Gly							
	50					55									

<210> 106
<211> 59
<212> PRT
<213> Dendroaspis angusticeps (Eastern green mamba C13 S2 C3 toxin)

<220>
<221> misc_feature
<222> (1)..(1)
<223> Xaa is Glu or Gln

<400> 106

Xaa	Pro	Arg	Arg	Lys	Leu	Cys	Ile	Leu	His	Arg	Asn	Pro	Gly	Arg	Cys
1				5					10					15	
Tyr	Asp	Lys	Ile	Pro	Ala	Phe	Tyr	Tyr	Asn	Gln	Lys	Lys	Lys	Gln	Cys
			20					25					30		
Glu	Arg	Phe	Asp	Trp	Ser	Gly	Cys	Gly	Gly	Asn	Ser	Asn	Arg	Phe	Lys
		35					40					45			
Thr	Ile	Glu	Glu	Cys	Arg	Arg	Thr	Cys	Ile	Gly					
	50					55									

<210> 107
<211> 57
<212> PRT
<213> Dendroaspis polylepis polylepis (Black mamba B toxin)

<400> 107

Arg	Pro	Tyr	Ala	Cys	Glu	Leu	Ile	Val	Ala	Ala	Gly	Pro	Cys	Met	Phe
1				5					10					15	
Phe	Ile	Ser	Ala	Phe	Tyr	Tyr	Ser	Lys	Gly	Ala	Asn	Lys	Cys	Tyr	Pro
			20					25					30		
Phe	Thr	Tyr	Ser	Gly	Cys	Arg	Gly	Asn	Ala	Asn	Arg	Phe	Lys	Thr	Ile
		35					40					45			

Glu Glu Cys Arg Arg Thr Cys Val Val
50 55

<210> 108
<211> 59
<212> PRT
<213> Dendroaspis polylepis polylepis (Black mamba E toxin)

<400> 108

Leu Gln His Arg Thr Phe Cys Lys Leu Pro Ala Glu Pro Gly Pro Cys
1 5 10 15

Lys Ala Ser Ile Pro Ala Phe Tyr Tyr Asn Trp Ala Ala Lys Lys Cys
20 25 30

Gln Leu Phe His Tyr Gly Gly Cys Lys Gly Asn Ala Asn Arg Phe Ser
35 40 45

Thr Ile Glu Lys Cys Arg His Ala Cys Val Gly
50 55

<210> 109
<211> 61
<212> PRT
<213> Vipera ammodytes TI toxin

<220>
<221> misc_feature
<222> (1)..(1)
<223> Xaa is Glu or Gln

<400> 109

Xaa Asp His Pro Lys Phe Cys Tyr Leu Pro Ala Asp Pro Gly Arg Cys
1 5 10 15

Lys Ala His Ile Pro Arg Phe Tyr Tyr Asp Ser Ala Ser Asn Lys Cys
20 25 30

Asn Lys Phe Ile Tyr Gly Gly Cys Pro Gly Asn Ala Asn Asn Phe Lys
35 40 45

Thr Trp Asp Glu Cys Arg Gln Thr Cys Gly Ala Ser Ala
50 55 60

<210> 110
<211> 62
<212> PRT
<213> Vipera ammodytes CTI toxin

<400> 110

Arg Asp Arg Pro Lys Phe Cys Tyr Leu Pro Ala Asp Pro Gly Arg Cys
1 5 10 15

Leu Ala Tyr Met Pro Arg Phe Tyr Tyr Asn Pro Ala Ser Asn Lys Cys
20 25 30

Glu Lys Phe Ile Tyr Gly Gly Cys Arg Gly Asn Ala Asn Asn Phe Lys
35 40 45

Thr Trp Asp Glu Cys Arg His Thr Cys Val Ala Ser Gly Ile
50 55 60

<210> 111

<211> 62

<212> PRT

<213> Bungarus fasciatus VIII B toxin

<400> 111

Lys Asn Arg Pro Thr Phe Cys Asn Leu Leu Pro Glu Thr Gly Arg Cys
1 5 10 15

Asn Ala Leu Ile Pro Ala Phe Tyr Tyr Asn Ser His Leu His Lys Cys
20 25 30

Gln Lys Phe Asn Tyr Gly Gly Cys Gly Gly Asn Ala Asn Asn Phe Lys
35 40 45

Thr Ile Asp Glu Cys Gln Arg Thr Cys Ala Ala Lys Tyr Gly
50 55 60

<210> 112

<211> 59

<212> PRT

<213> Anemonia sulcata

<400> 112

Ile Asn Gly Asp Cys Glu Leu Pro Lys Val Val Gly Pro Cys Arg Ala
1 5 10 15

Arg Phe Pro Arg Tyr Tyr Tyr Asn Ser Ser Ser Lys Arg Cys Glu Lys
20 25 30

Phe Ile Tyr Gly Gly Cys Gly Gly Asn Ala Asn Asn Phe His Thr Leu
35 40 45

Glu Glu Cys Glu Lys Val Cys Gly Val Arg Ser
50 55

<210> 113

<211> 56

<212> PRT

<213> Homo sapiens

<400> 113

Lys Glu Asp Ser Cys Gln Leu Gly Tyr Ser Ala Gly Pro Cys Met Gly
1 5 10 15

Met Thr Ser Arg Tyr Phe Tyr Asn Gly Thr Ser Met Ala Cys Glu Thr
20 25 30

Phe Gln Tyr Gly Gly Cys Met Gly Asn Gly Asn Asn Phe Val Thr Glu
35 40 45

Lys Glu Cys Leu Gln Thr Cys Arg
50 55

<210> 114
<211> 61
<212> PRT
<213> Homo sapiens

<400> 114

Thr Val Ala Ala Cys Asn Leu Pro Val Ile Arg Gly Pro Cys Arg Ala
1 5 10 15

Phe Ile Gln Leu Trp Ala Phe Asp Ala Val Lys Gly Lys Cys Val Leu
20 25 30

Phe Pro Tyr Gly Gly Cys Gln Gly Asn Gly Asn Lys Phe Tyr Ser Glu
35 40 45

Lys Glu Cys Arg Glu Tyr Cys Gly Val Pro Gly Asp Glu
50 55 60

<210> 115
<211> 60
<212> PRT
<213> Bungarus multicinctus (beta bungarotoxin B1)

<400> 115

Arg Gln Arg His Arg Asp Cys Asp Lys Pro Pro Asp Lys Gly Asn Cys
1 5 10 15

Gly Pro Val Arg Ala Phe Tyr Tyr Asp Thr Arg Leu Lys Thr Cys Lys
20 25 30

Ala Phe Gln Tyr Arg Gly Cys Asp Gly Asp His Gly Asn Phe Lys Thr
35 40 45

Glu Thr Leu Cys Arg Cys Glu Cys Leu Val Tyr Pro
50 55 60

<210> 116
<211> 60
<212> PRT
<213> Bungarus multicinctus (beta bungarotoxin B2)

<400> 116

Arg Lys Arg His Pro Asp Cys Asp Lys Pro Pro Asp Thr Lys Ile Cys
1 5 10 15

Gln Thr Val Arg Ala Phe Tyr Tyr Lys Pro Ser Ala Lys Arg Cys Val
20 25 30

Gln Phe Arg Tyr Gly Gly Cys Asp Gly Asp His Gly Asn Phe Lys Ser
35 40 45

Asp His Leu Cys Arg Cys Glu Cys Glu Leu Tyr Arg
50 55 60

<210> 117
<211> 58
<212> PRT
<213> Bos taurus

<400> 117

Arg Pro Asp Phe Cys Leu Glu Pro Pro Tyr Thr Gly Pro Cys Lys Ala
1 5 10 15

Lys Met Ile Arg Tyr Phe Tyr Asn Ala Lys Ala Gly Phe Cys Glu Thr
20 25 30

Phe Val Tyr Gly Gly Cys Lys Ala Lys Ser Asn Asn Phe Arg Ser Ala
35 40 45

Glu Asp Cys Met Arg Thr Cys Gly Gly Ala
50 55

<210> 118
<211> 61
<212> PRT
<213> Tachypleus tridentatus

<400> 118

Thr Glu Arg Gly Phe Leu Asp Cys Thr Ser Pro Pro Val Thr Gly Pro
1 5 10 15

Cys Arg Ala Gly Phe Lys Arg Tyr Asn Tyr Asn Thr Arg Thr Lys Gln
20 25 30

Cys Glu Pro Phe Lys Tyr Gly Gly Cys Lys Gly Asn Gly Asn Arg Tyr
35 40 45

Lys Ser Glu Gln Asp Cys Leu Asp Ala Cys Ser Gly Phe
50 55 60

<210> 119
<211> 62
<212> PRT
<213> Bombyx mori

<220>
<221> misc_feature
<222> (14)..(14)
<223> Xaa is Phe or Gly

<400> 119

Asp Glu Pro Thr Thr Asp Leu Pro Ile Cys Glu Gln Ala Xaa Asp Ala
1 5 10 15

Gly Leu Cys Phe Gly Tyr Met Lys Leu Tyr Ser Tyr Asn Gln Glu Thr
20 25 30

Lys Asn Cys Glu Glu Phe Ile Tyr Gly Gly Cys Gln Gly Asn Asp Asn
35 40 45

Arg Phe Ser Thr Leu Ala Glu Cys Glu Gln Lys Cys Ile Asn
50 55 60

<210> 120

<211> 56

<212> PRT

<213> Bos taurus

<400> 120

Lys Ala Asp Ser Cys Gln Leu Asp Tyr Ser Gln Gly Pro Cys Leu Gly
1 5 10 15

Leu Phe Lys Arg Tyr Phe Tyr Asn Gly Thr Ser Met Ala Cys Glu Thr
20 25 30

Phe Leu Tyr Gly Gly Cys Met Gly Asn Leu Asn Asn Phe Leu Ser Gln
35 40 45

Lys Glu Cys Leu Gln Thr Cys Arg
50 55

<210> 121

<211> 61

<212> PRT

<213> Bos taurus

<400> 121

Thr Val Glu Ala Cys Asn Leu Pro Ile Val Gln Gly Pro Cys Arg Ala
1 5 10 15

Phe Ile Gln Leu Trp Ala Phe Asp Ala Val Lys Gly Lys Cys Val Arg
20 25 30

Phe Ser Tyr Gly Gly Cys Lys Gly Asn Gly Asn Lys Phe Tyr Ser Gln
35 40 45

Lys Glu Cys Lys Glu Tyr Cys Gly Ile Pro Gly Glu Ala
50 55 60

<210> 122

<211> 58

<212> PRT

<213> Artificial Sequence

<220>

<223> Engineered BPTI (KR15, ME52)

<400> 122

Arg Pro Asp Phe Cys Leu Glu Pro Pro Tyr Thr Gly Pro Cys Arg Ala
1 5 10 15
Arg Ile Ile Arg Tyr Phe Tyr Asn Ala Lys Ala Gly Leu Cys Gln Thr
20 25 30
Phe Val Tyr Gly Gly Cys Arg Ala Lys Arg Asn Asn Phe Lys Ser Ala
35 40 45
Glu Asp Cys Glu Arg Thr Cys Gly Gly Ala
50 55

<210> 123

<211> 59

<212> PRT

<213> Artificial Sequence

<220>

<223> Isoaprotinin G-1

<220>

<221> misc_feature

<222> (1)..(1)

<223> Xaa is Glu or Gln

<400> 123

Xaa Arg Pro Asp Phe Cys Leu Glu Pro Pro Tyr Thr Gly Pro Cys Lys
1 5 10 15
Ala Arg Met Ile Arg Tyr Phe Tyr Asn Ala Lys Ala Gly Leu Cys Gln
20 25 30
Pro Phe Val Tyr Gly Gly Cys Arg Ala Lys Ser Asn Asn Phe Lys Ser
35 40 45
Ala Glu Asp Cys Met Arg Thr Cys Gly Gly Ala
50 55

<210> 124

<211> 58

<212> PRT

<213> Artificial Sequence

<220>

<223> Isoaprotinin 2

<400> 124

Arg Pro Asp Phe Cys Leu Glu Pro Pro Tyr Thr Gly Pro Cys Lys Ala
1 5 10 15
Arg Ile Ile Arg Tyr Phe Tyr Asn Ala Lys Ala Gly Leu Cys Gln Pro
20 25 30

Phe Val Tyr Gly Gly Cys Arg Ala Lys Arg Asn Asn Phe Lys Ser Ser
35 40 45

Glu Asp Cys Met Arg Thr Cys Gly Gly Ala
50 55

<210> 125
<211> 58
<212> PRT
<213> Artificial Sequence

<220>

<223> Isoaprotinin G-2

<400> 125

Arg Pro Asp Phe Cys Leu Glu Pro Pro Tyr Thr Gly Pro Cys Lys Ala
1 5 10 15

Arg Met Ile Arg Tyr Phe Tyr Asn Ala Lys Ala Gly Leu Cys Gln Pro
20 25 30

Phe Val Tyr Gly Gly Cys Arg Ala Lys Arg Asn Asn Phe Lys Ser Ala
35 40 45

Glu Asp Cys Met Arg Thr Cys Gly Gly Ala
50 55

<210> 126
<211> 58
<212> PRT
<213> Artificial Sequence

<220>

<223> Isoaprotinin 1

<400> 126

Arg Pro Asp Phe Cys Leu Glu Pro Pro Tyr Thr Gly Pro Cys Lys Ala
1 5 10 15

Lys Met Ile Arg Tyr Phe Tyr Asn Ala Lys Ala Gly Phe Cys Glu Thr
20 25 30

Phe Val Tyr Gly Gly Cys Lys Ala Lys Ser Asn Asn Phe Arg Ser Ala
35 40 45

Glu Asp Cys Met Arg Thr Cys Gly Gly Ala
50 55

<210> 127
<211> 11
<212> DNA
<213> Artificial Sequence

<220>

<223> PflMI restriction site

<220>

<221> misc_feature

<222> (4)..(8)

<223> n is a, c, g or t

<400> 127

ccannnnntg g

11

<210> 128

<211> 15

<212> DNA

<213> Artificial Sequence

<220>

<223> XcmI restriction site

<220>

<221> misc_feature

<222> (4)..(12)

<223> n is a, c, g or t

<400> 128

ccannnnnnn nntgg

15

<210> 129

<211> 9

<212> PRT

<213> Artificial Sequence

<220>

<223> amino acids 13-21 of EpiNE alpha

<400> 129

Pro Cys Val Ala Met Phe Gln Arg Tyr

1

5

<210> 130

<211> 6

<212> PRT

<213> Artificial Sequence

<220>

<223> amino acids 15-20 of EpiNE-7

<400> 130

Val Ala Met Phe Pro Arg

1

5

<210> 131
<211> 4
<212> PRT
<213> Artificial Sequence

<220>

<223> amino acids 35-38 of HNE

<400> 131

Tyr Gly Gly Cys
1

<210> 132
<211> 9
<212> PRT
<213> Artificial Sequence

<220>

<223> amino acids 13-21 of BPTI

<400> 132

Pro Cys Lys Ala Arg Ile Ile Arg Tyr
1 5

<210> 133
<211> 9
<212> PRT
<213> Artificial Sequence

<220>

<223> amino acids 13-21 of EpiNE3

<400> 133

Pro Cys Val Gly Phe Phe Ser Arg Tyr
1 5

<210> 134
<211> 9
<212> PRT
<213> Artificial Sequence

<220>

<223> amino acids 13-21 of EpiNE6

<400> 134

Pro Cys Val Gly Phe Phe Gln Arg Tyr
1 5

<210> 135
<211> 9
<212> PRT
<213> Artificial Sequence

<220>

<223> amino acids 13-21 of EpiNE7

<400> 135

Pro Cys Val Ala Met Phe Pro Arg Tyr
1 5

<210> 136
<211> 9
<212> PRT
<213> Artificial Sequence

<220>

<223> amino acids 13-21 of EpiNE4

<400> 136

Pro Cys Val Ala Ile Phe Pro Arg Tyr
1 5

<210> 137
<211> 9
<212> PRT
<213> Artificial Sequence

<220>

<223> amino acids 13-21 of EpiNE8

<400> 137

Pro Cys Val Ala Ile Phe Lys Arg Ser
1 5

<210> 138
<211> 9
<212> PRT
<213> Artificial Sequence

<220>

<223> amino acids 13-21 of EpiNE1

<400> 138

Pro Cys Ile Ala Phe Phe Pro Arg Tyr
1 5

<210> 139
<211> 9
<212> PRT
<213> Artificial Sequence

<220>

<223> amino acids 13-21 of EpiNE5

<400> 139

Pro Cys Ile Ala Phe Phe Gln Arg Tyr
1 5

<210> 140
<211> 9
<212> PRT
<213> Artificial Sequence

<220>

<223> amino acids 13-21 of EpiNE2

<400> 140

Pro Cys Ile Ala Leu Phe Lys Arg Tyr
1 5